

# Lecture 6

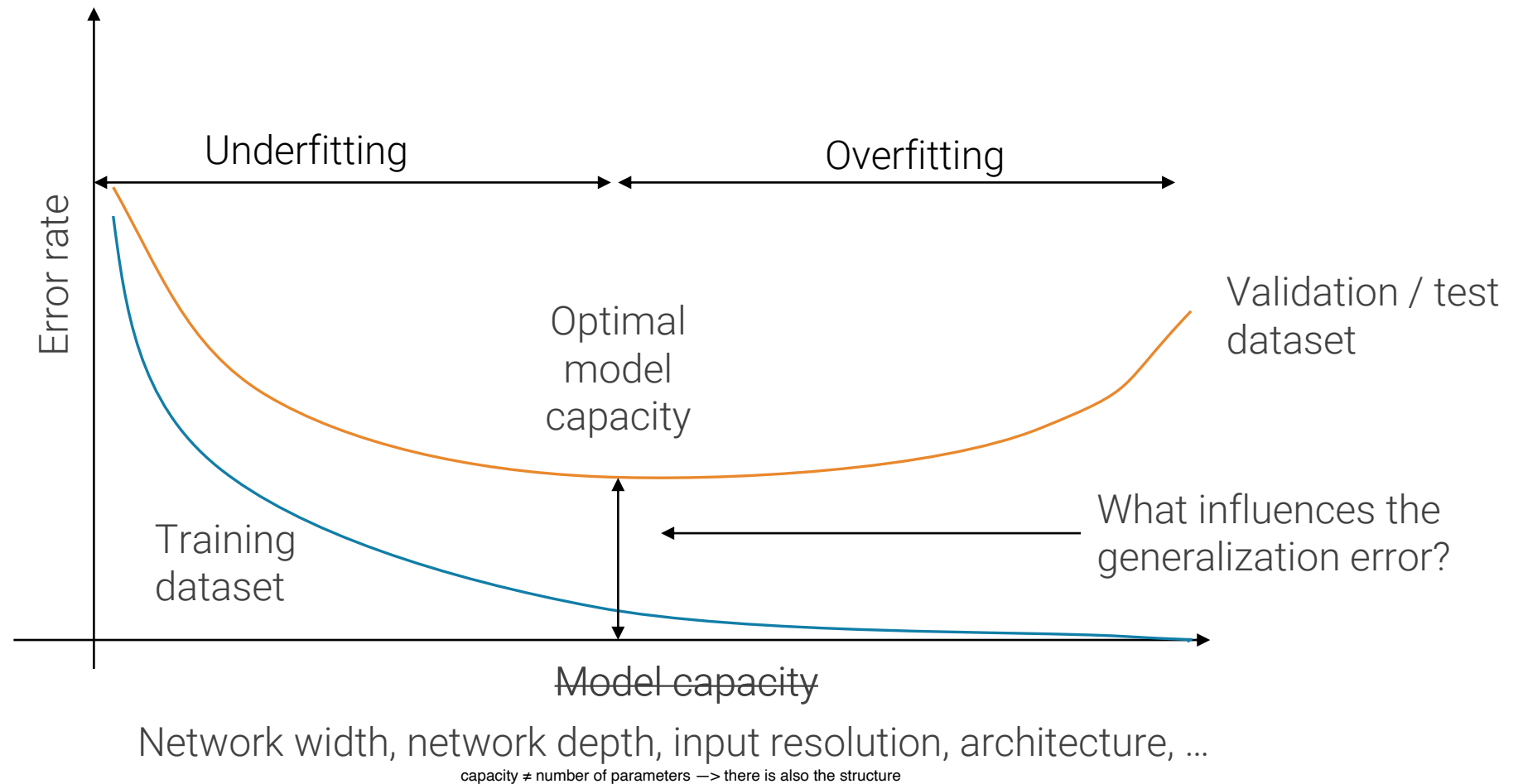
## Regularization and training recipes

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IMAGE PROCESSING AND COMPUTER VISION – PART 2

SAMUELE SALTI

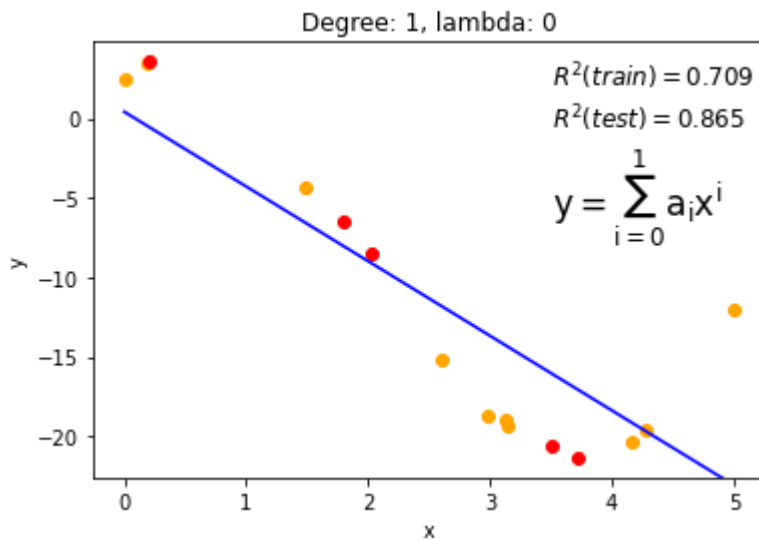
# Generalization error



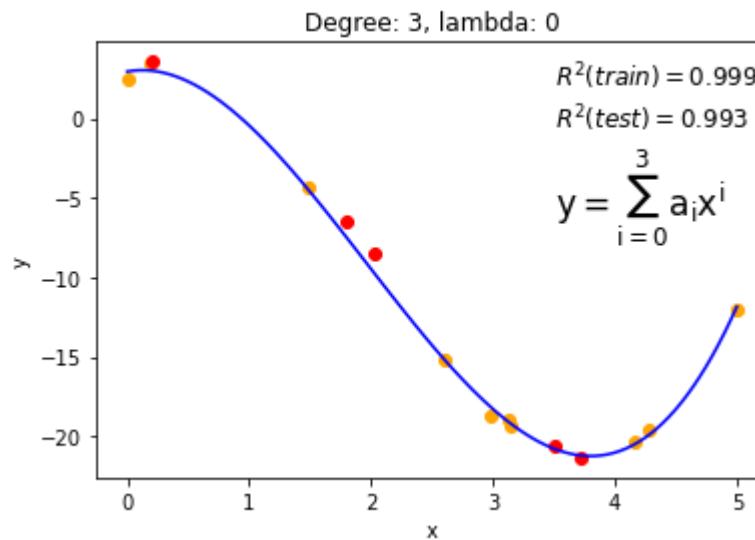
# Example: generalized linear regression

Suppose we fit  $N$  data points, generated by a noisy cubic equation  $y = (ax^3 + bx^2 + cx + d) + \epsilon$  with different models, which are polynomials of degree  $K$  of the form  $y = \sum_{i=0}^K a_i x^i = a_0 + a_1x + a_2x^2 + a_3x^3 + \dots$

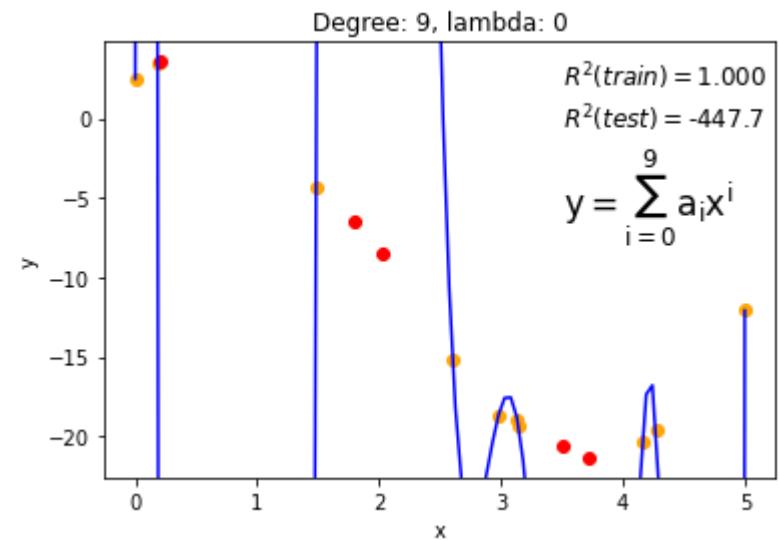
Linear model  
Underfitting



“Right”  
model



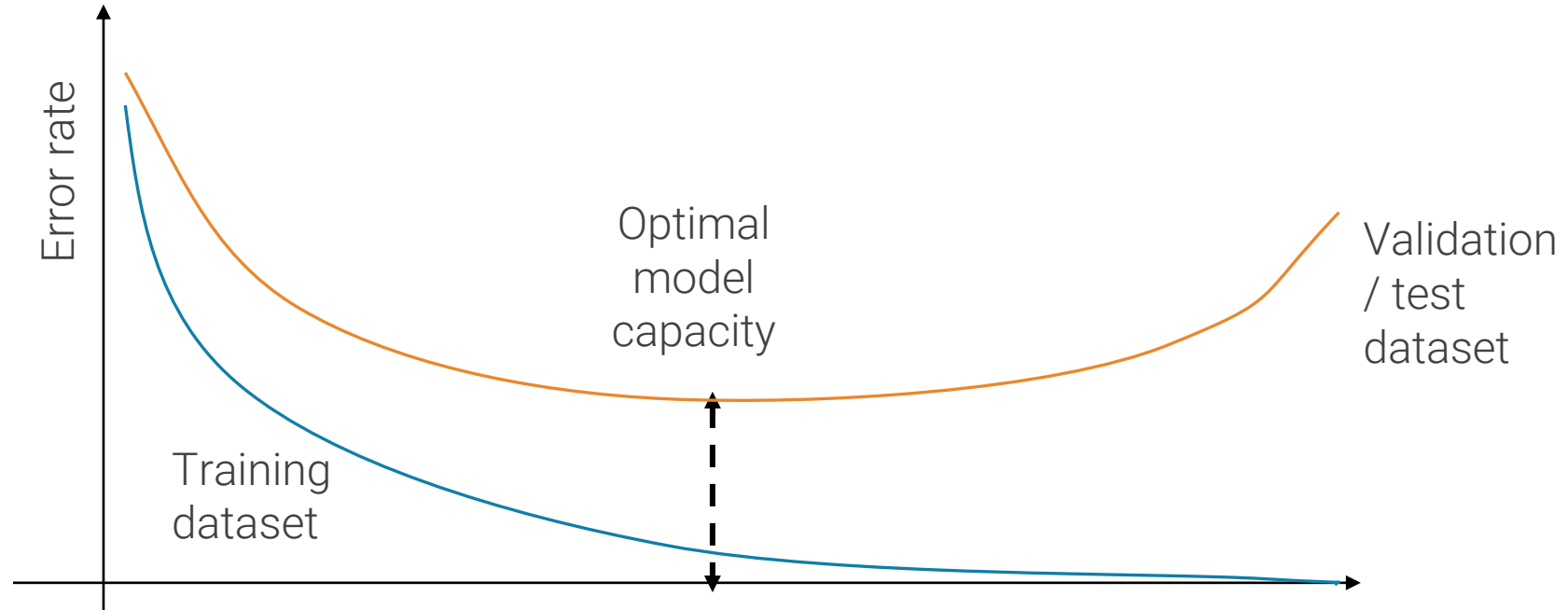
“Interpolating” model  
Overfitting



<http://www.deeplearningbook.org/contents/ml.html>

# Effective capacity

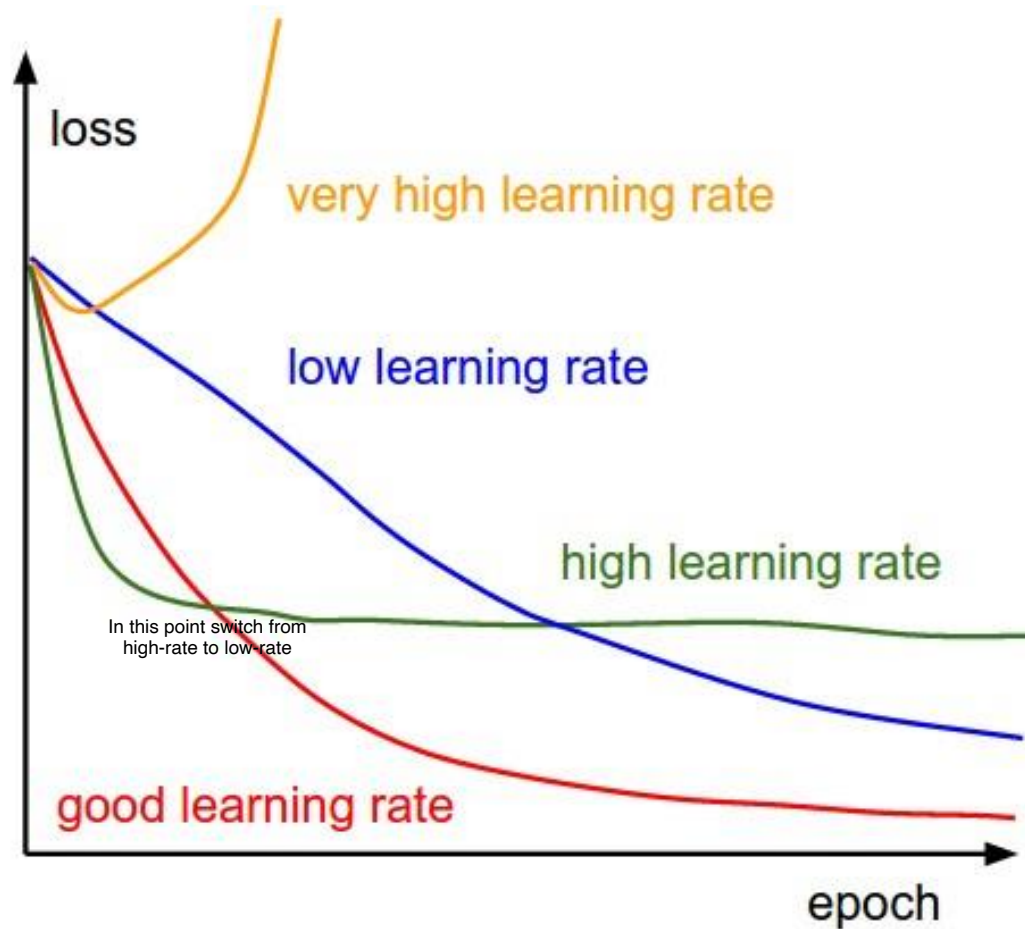
In practice, we very often train a specific architecture, whose theoretical capacity is fixed and large. Yet, the **effective capacity** of an architecture can still be changed due to **optimization hyperparameters** and **regularization**.



Theoretical capacity: network width, network depth, input resolution, etc..

Effective capacity: learning rate, favor small values for parameters, training time, ...

# Learning rate is a key hyperparameter

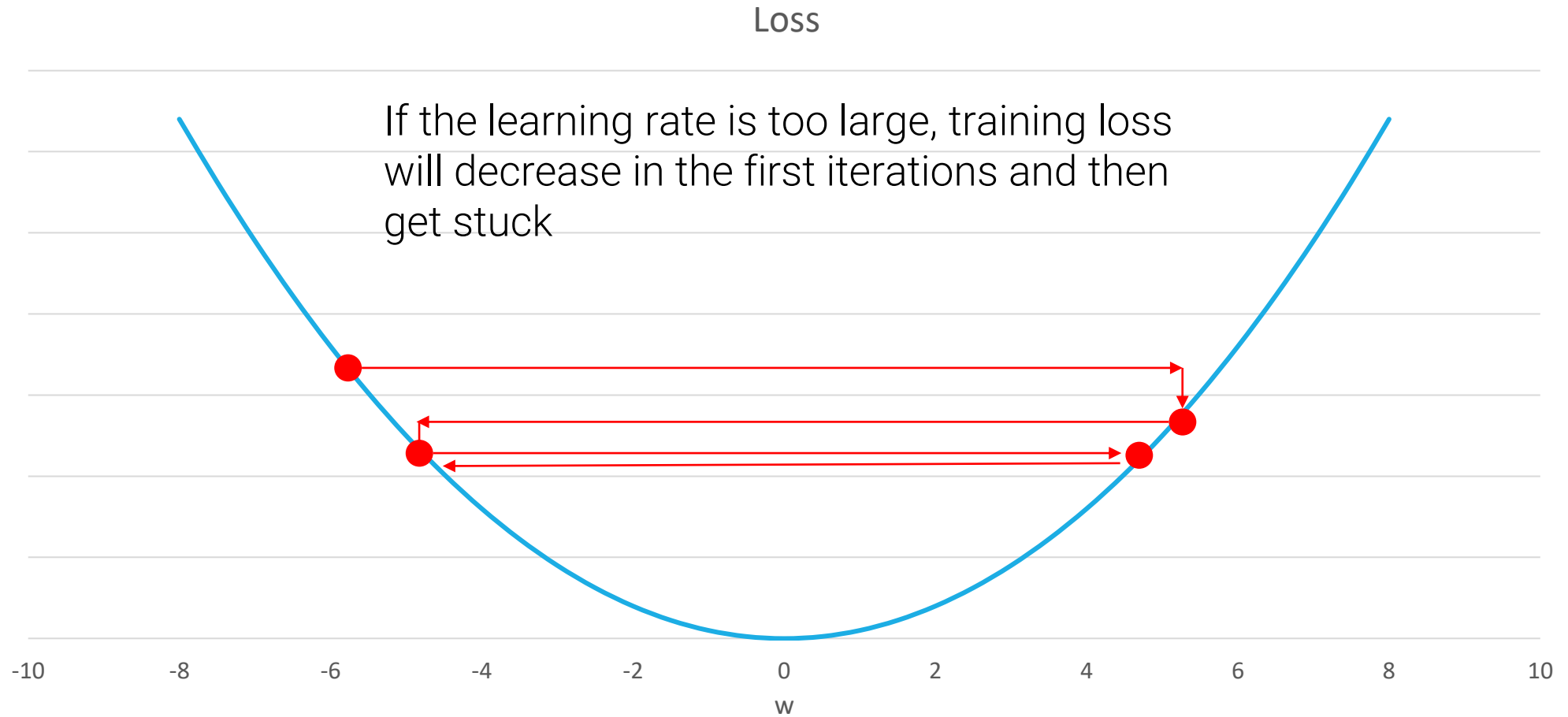


Problem: hard to find the good learning rate to reach good fitting

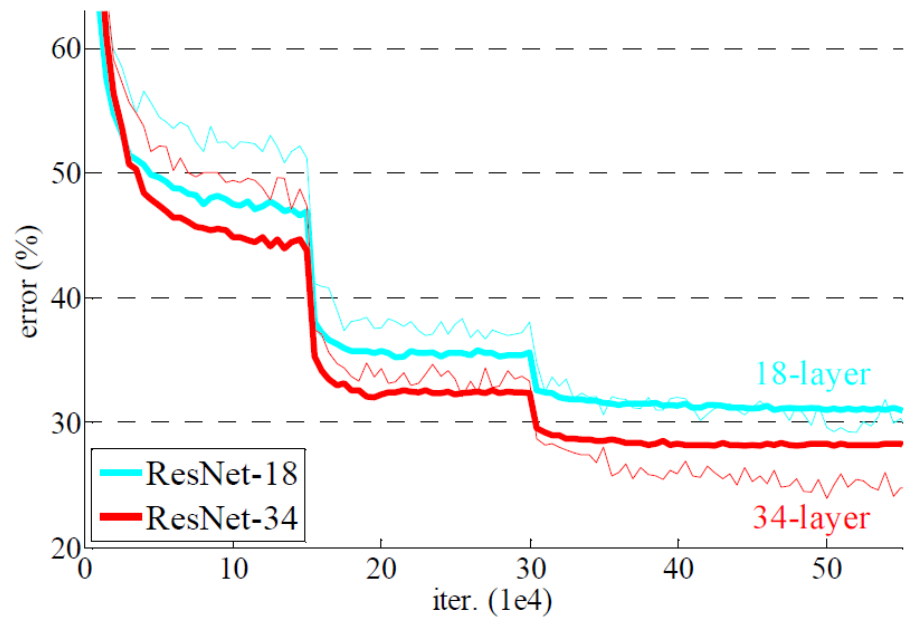
Solution: use a mixture of high and low.

Source: <https://cs231n.github.io/neural-networks-3/>

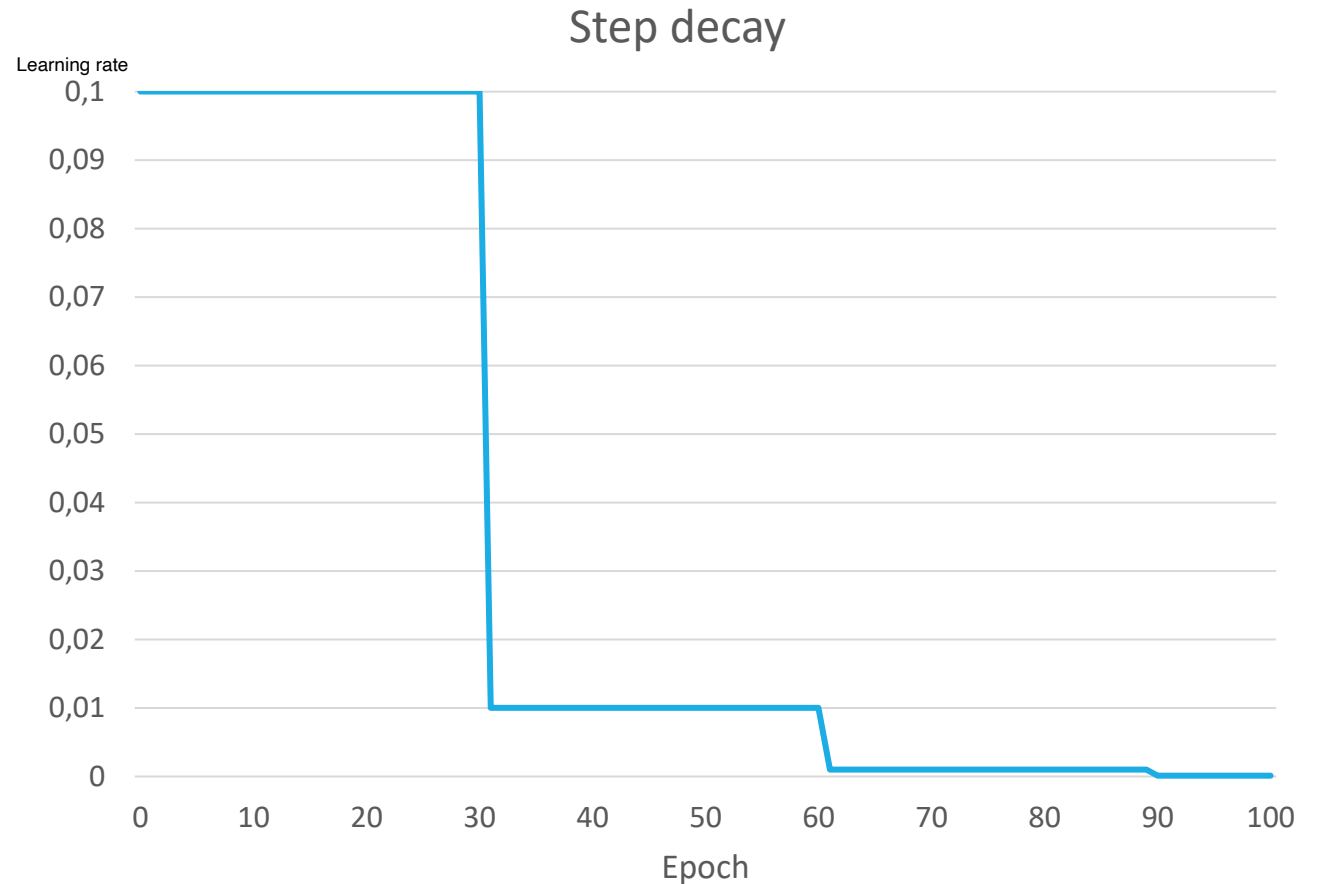
# Learning rate and gradient descent



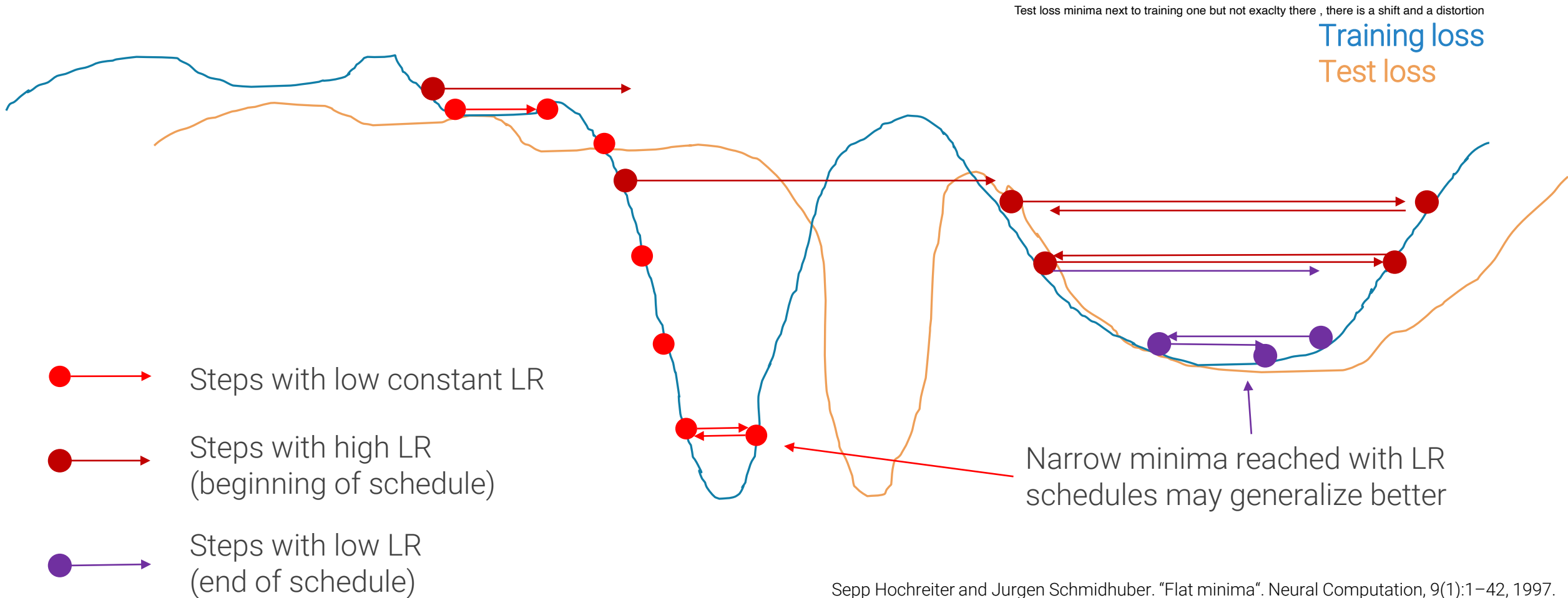
# Learning rate schedule: step



Step schedule (or decay): start with high learning rate (e.g. 0.1) and divide by 10 when the error plateaus  
Used e.g., in ResNets



# Intuition: favor wide minima



Sepp Hochreiter and Jurgen Schmidhuber. "Flat minima". Neural Computation, 9(1):1–42, 1997.

Laurent Dinh et al., "Sharp minima can generalize for deep nets.", ICLR 2017.

Nitish Shirish Keskar et al., "On large-batch training for deep learning: Generalization gap and sharp minima." ICLR 2017.

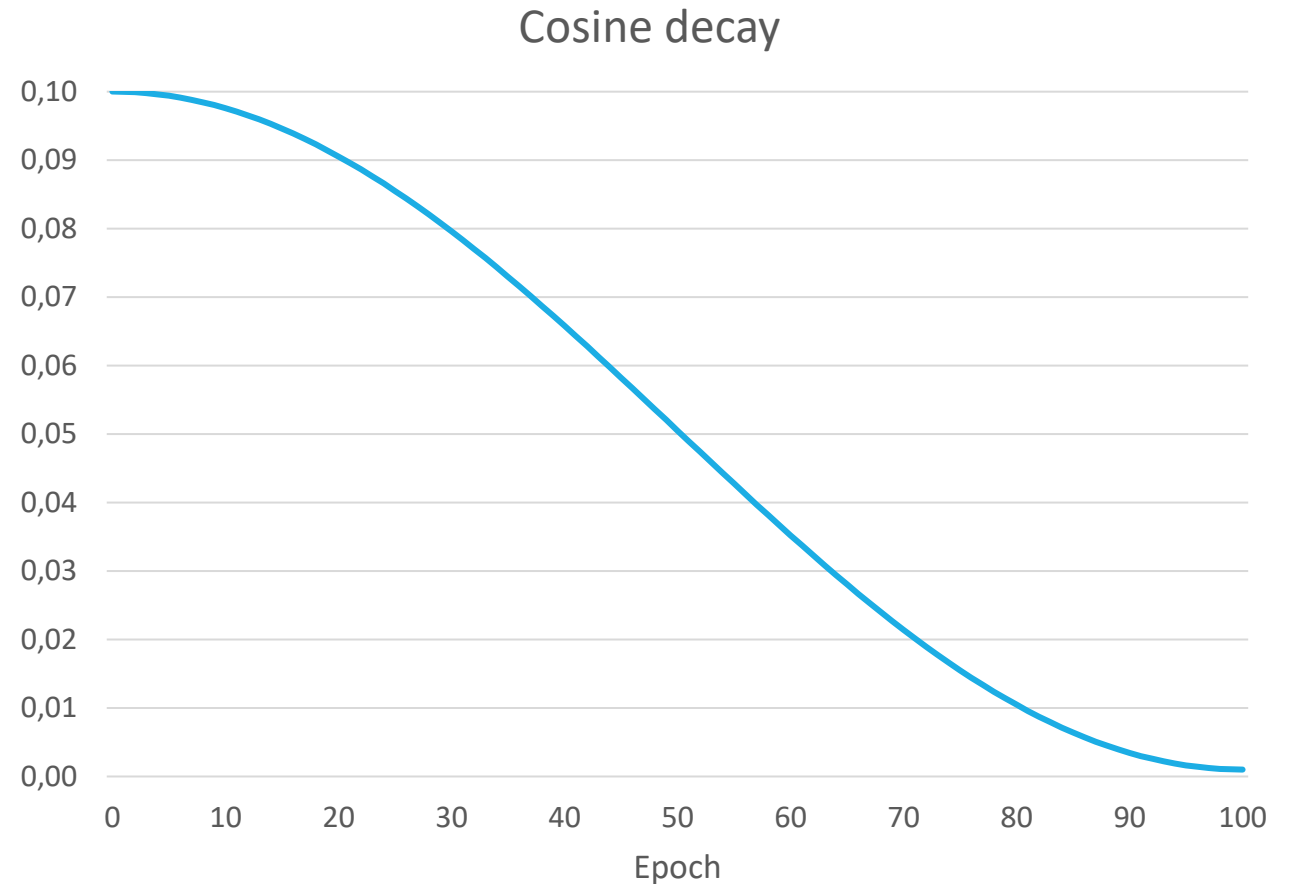


# Learning rate schedule: cosine

Cosine schedule (or decay): if training for  $E$  epochs, learning rate for epoch  $e$  is

$$lr_e = lr_E + \frac{1}{2}(lr_0 - lr_E) \left( 1 + \cos\left(\frac{e \pi}{E}\right) \right)$$

It lets the optimizer follow a similar path to step decays with less hyper parameters to tune.



Ilya Loshchilov, Frank Hutter "SGDR: Stochastic Gradient Descent with Warm Restarts", ICLR 2017

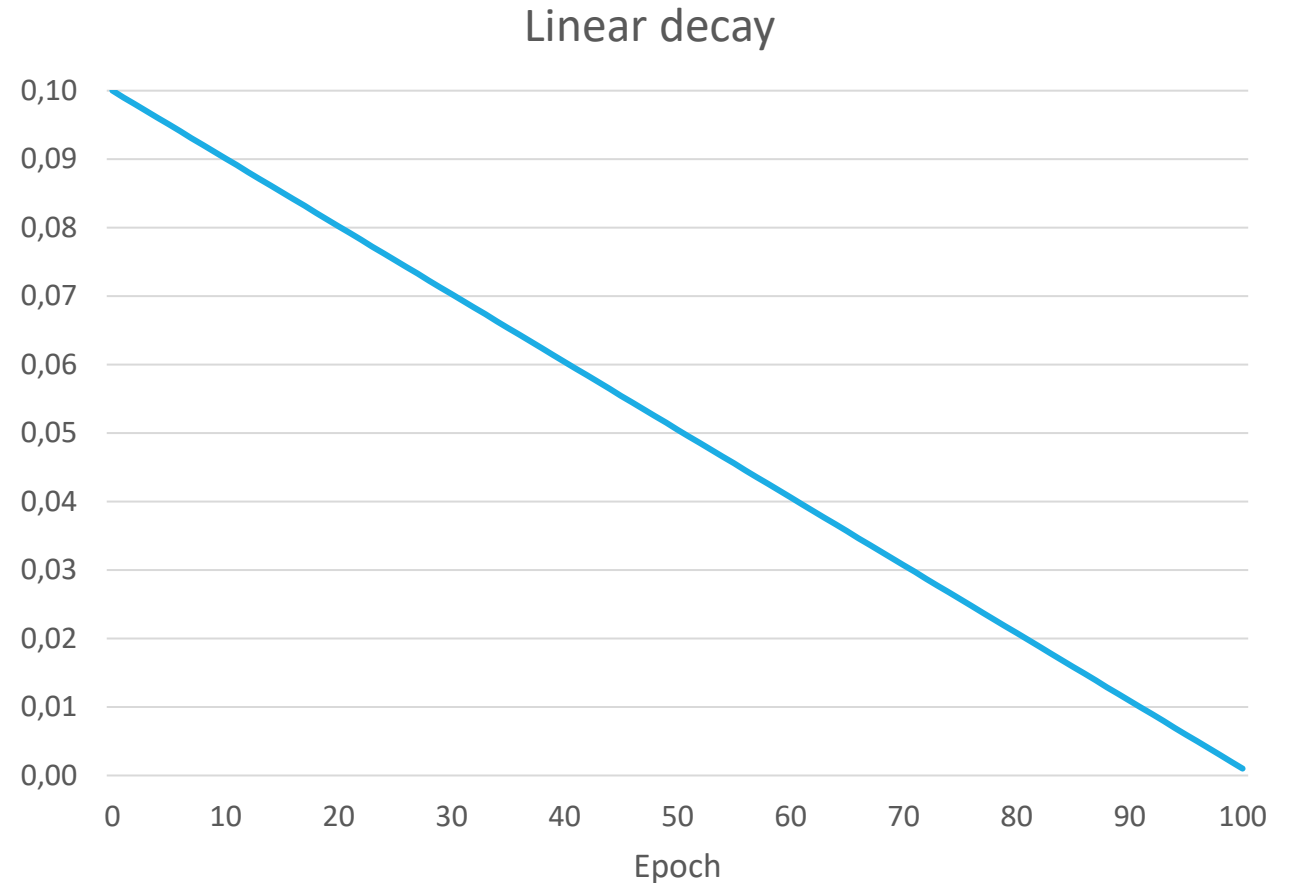
# Learning rate schedule: linear

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Linear schedule (or decay): if training for  $E$  epochs, learning rate for epoch  $e$  is

$$lr_e = lr_E + (lr_0 - lr_E) \left(1 - \frac{e}{E}\right)$$

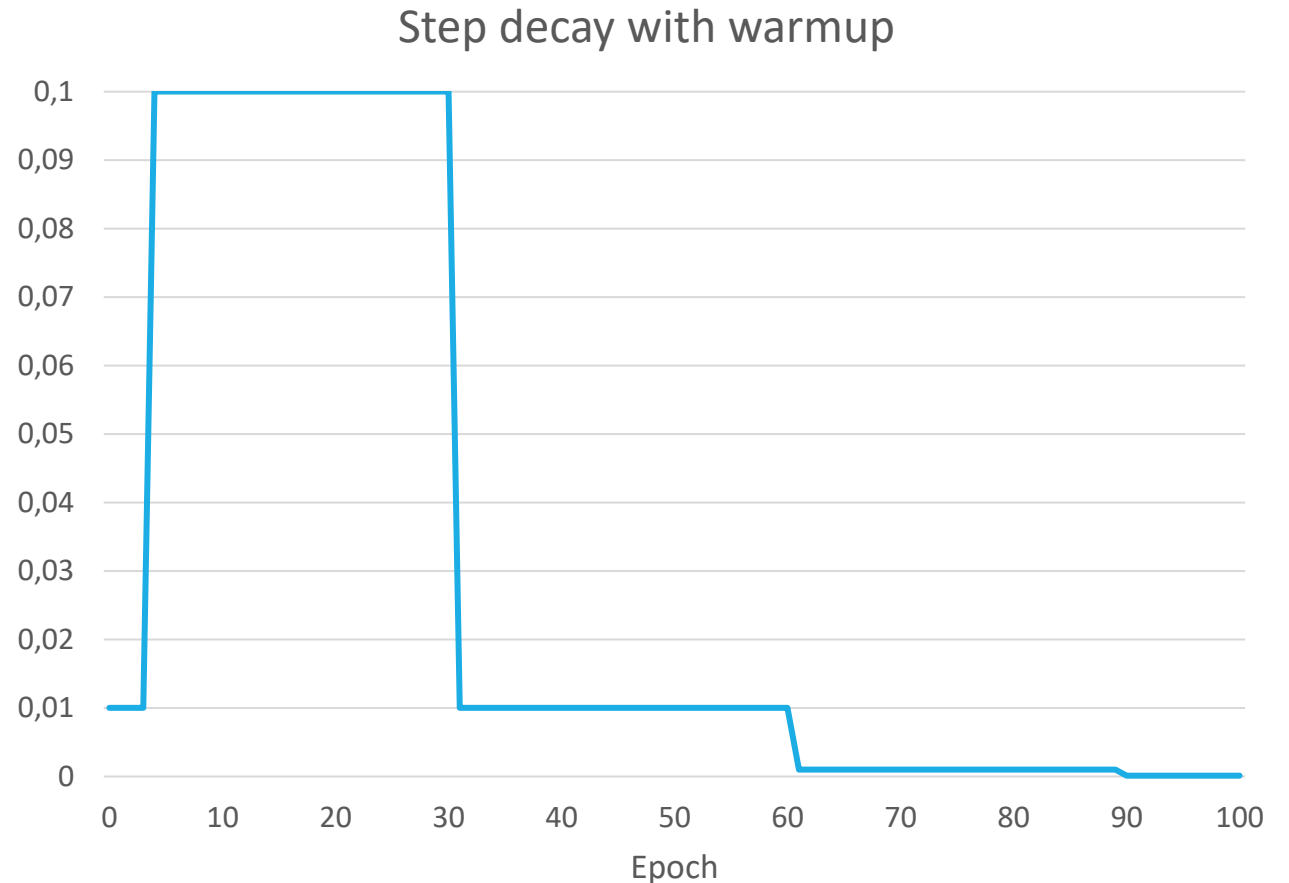
An even simpler alternative to coarsely emulate step decay evolution



# Learning rate schedule: warm-up

Schedules usually start with high learning rates. For very deep networks (e.g. ResNet-110 on CIFAR-10) a high learning rate can slow down convergence at the beginning of training (i.e. accuracy remains at chance levels for several epochs). **This is usually a symptom of poor initialization:** a way to counteract it is to **use a lower learning rate for a few epochs** (even one or less, e.g. until accuracy increases).

Usually beneficial also for shallower, randomly initialized networks, as at the beginning of training we may be in a hard-to-navigate part of the loss landscape.



Kaiming He et al., "Deep Residual Learning for Image Recognition", CVPR 2016

# Learning rate schedule: One cycle

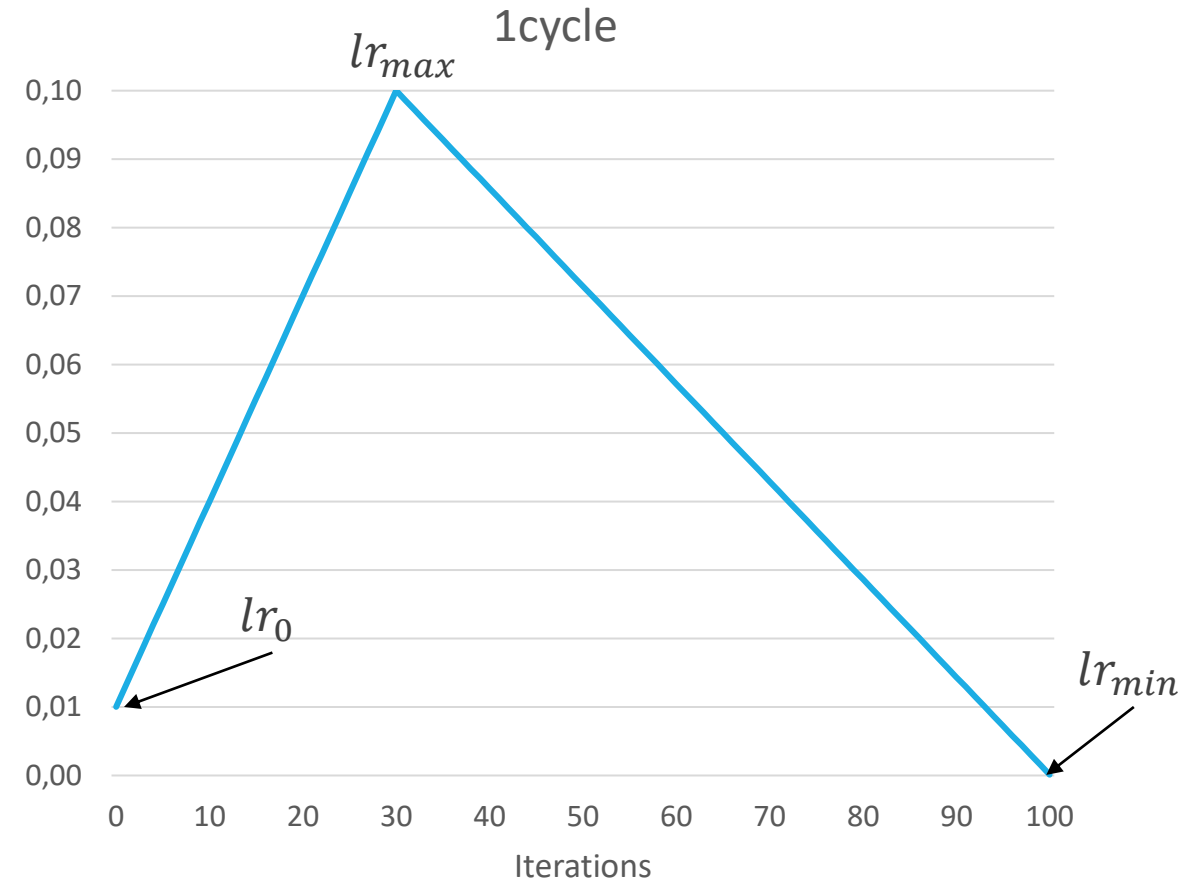
One cycle schedule (or decay) **modifies the learning rate after each mini-batch** (also referred to as iteration or step).

If training for a total number of iterations  $I$ , learning rate for each iteration  $i$  is

$$lr_i = \begin{cases} lr_{max} + (lr_0 - lr_{max}) \left(1 - \frac{i}{pI}\right) & \text{if } i < pI \\ lr_{min} + (lr_0 - lr_{min}) \left(1 - \frac{i - pI}{I - pI}\right) & \text{if } i \geq pI \end{cases}$$

where  $p$  is the fraction of iterations where we increase the learning rate (e.g. 0.3).

Original proposal had 3 phases, but PyTorch and other popular implementations provide two. It is usually realized with **cosine segments**.



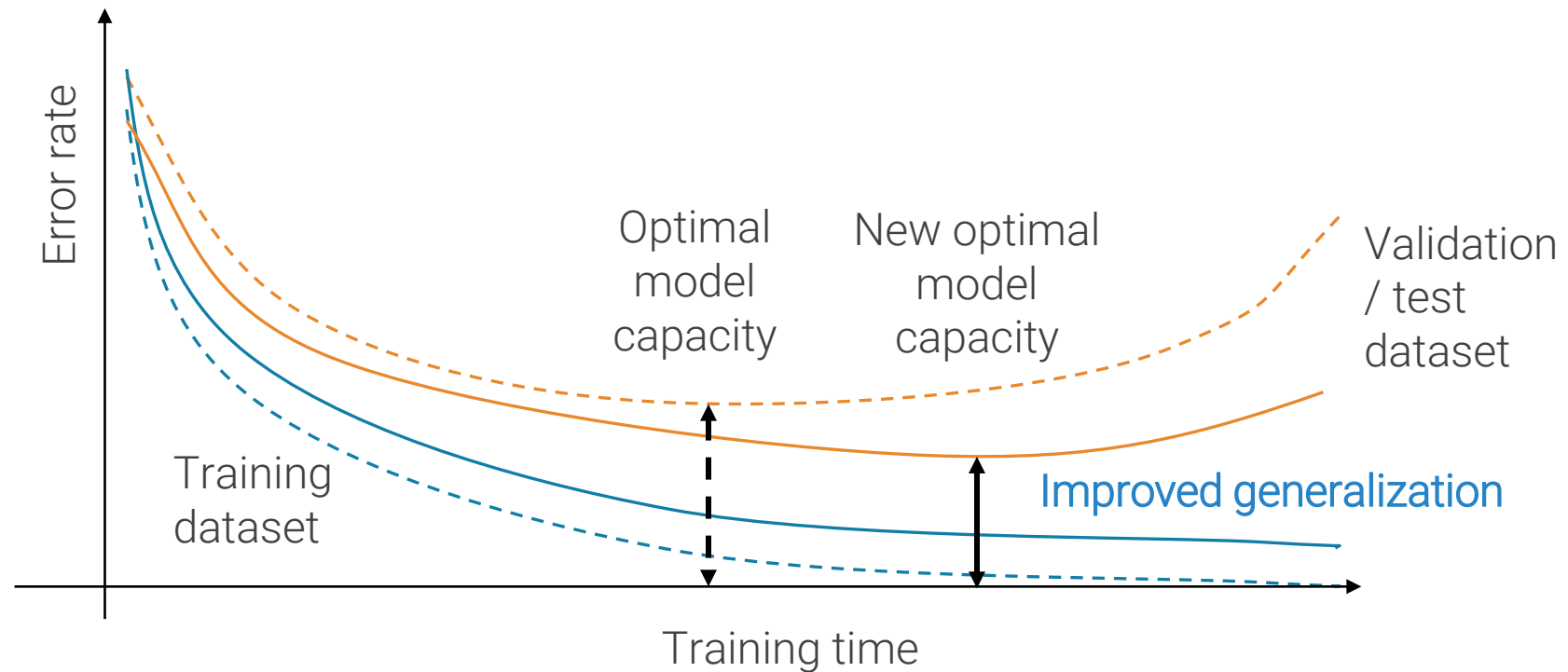
# Regularization

**Regularization** is any modification we make to the training recipe that is intended to

- reduce its generalization (test) error
- but **not** its training error.

We will review:

- parameter norm penalties
- early stopping
- label smoothing
- dropout
- stochastic depth
- data augmentation



# Parameter norm penalties

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When using parameter norm penalties, we (implicitly!) add a term to the loss

$$L(\theta; D^{train}) = \underbrace{L^{task}(\theta; D^{train})}_{\text{Task (or data) loss}} + \underbrace{\lambda L^{reg}(\theta)}_{\text{Regularization loss}}$$

**Task (or data) loss:** let training match the model to the input data and label, e.g. cross-entropy loss

**Regularization loss:** guide training to prefer “simpler” models

Commonly used norms are:

**$l_2$  regularization**  $L^{reg}(\theta) = \sum_i \theta_i^2$

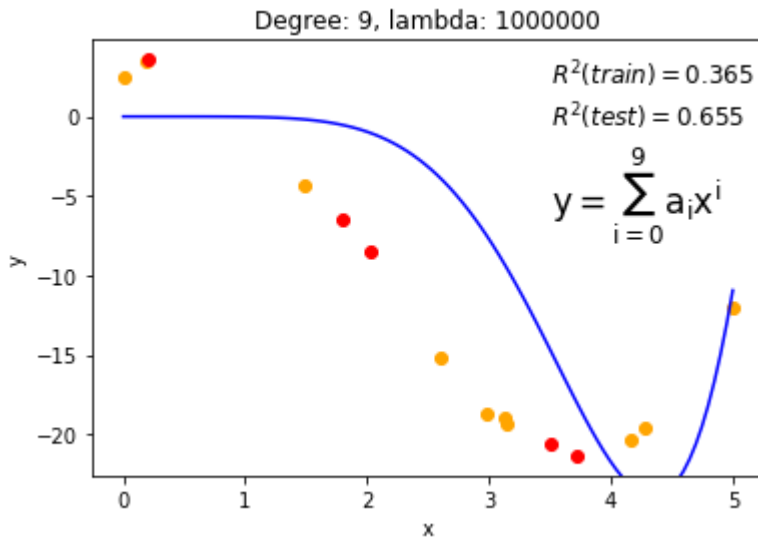
**$l_1$  regularization**  $L^{reg}(\theta) = \sum_i |\theta_i|$

Intuition: even if a model has millions of parameters, a constraint on the overall norm of the parameters forces them to be small, thereby limiting the **effective capacity** of my model

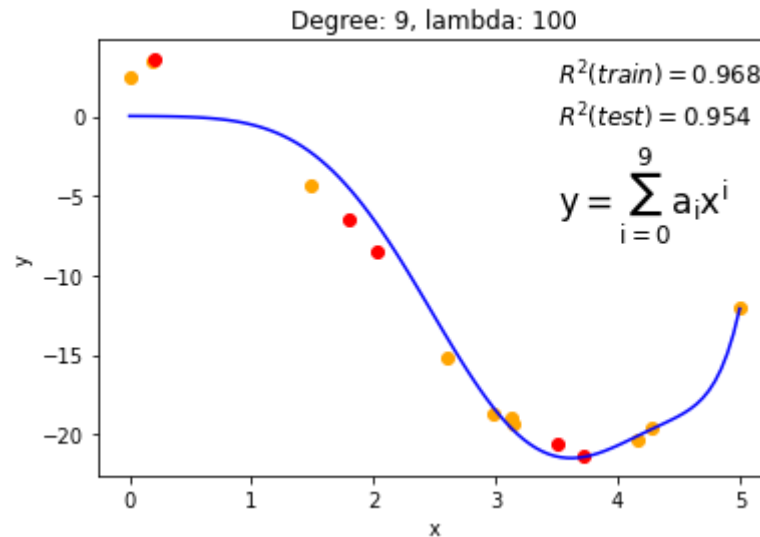
# Example: generalized linear regression with $l_2$ norm

We can fit the same  $N$  data points presented earlier with **high-capacity models**, i.e. polynomials of degree  $N$ , and control the ability of the model to generalize through appropriate regularization.

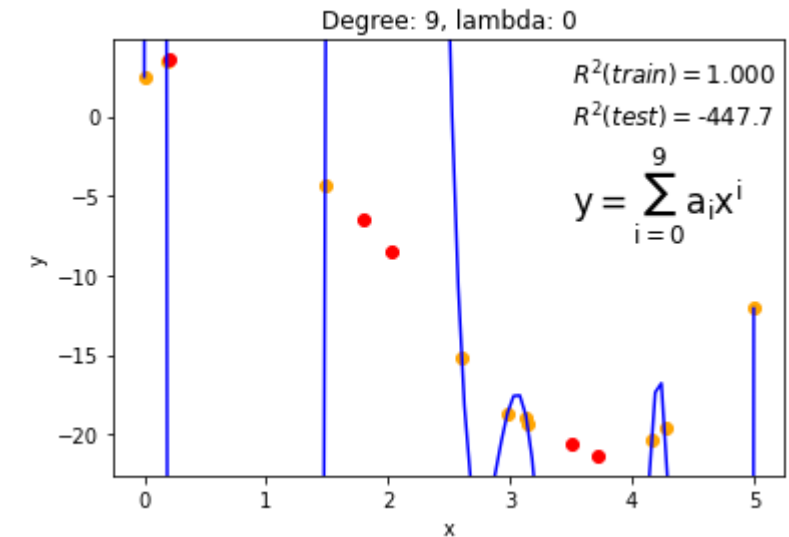
“Linear” model  
 $\lambda \rightarrow +\infty$



“Right” model  
appropriate  $\lambda$



“Interpolating” model  
 $\lambda \rightarrow 0$



<http://www.deeplearningbook.org/contents/ml.html>

# $l_2$ regularization or weight decay

$$L(\theta; D^{train}) = L^{task}(\theta; D^{train}) + \frac{\lambda}{2} \|\theta\|_2^2$$

$l_2$  regularization is also known as ridge regression or Tikhonov regularization

In deep learning, it also goes under the name of **weight decay** because **for plain SGD** every gradient descent step drives the weights toward the origin before applying the update used when regularization is not used

$$\begin{aligned}\theta^{(i+1)} &= \theta^{(i)} - lr \nabla_{\theta} L(\theta^{(i)}; D^{train}) \\ &= \theta^{(i)} - lr \nabla_{\theta} \left[ L^{task}(\theta^{(i)}; D^{train}) + \frac{\lambda}{2} \|\theta^{(i)}\|_2^2 \right] \\ &= \theta^{(i)} - lr \left[ \nabla_{\theta} L^{task}(\theta^{(i)}; D^{train}) + \lambda \theta^{(i)} \right]\end{aligned}$$

Decayed parameter vector	$= (1 - lr \lambda) \theta^{(i)} - lr \nabla_{\theta} L^{task}(\theta^{(i)}; D^{train})$	Gradient without regularization
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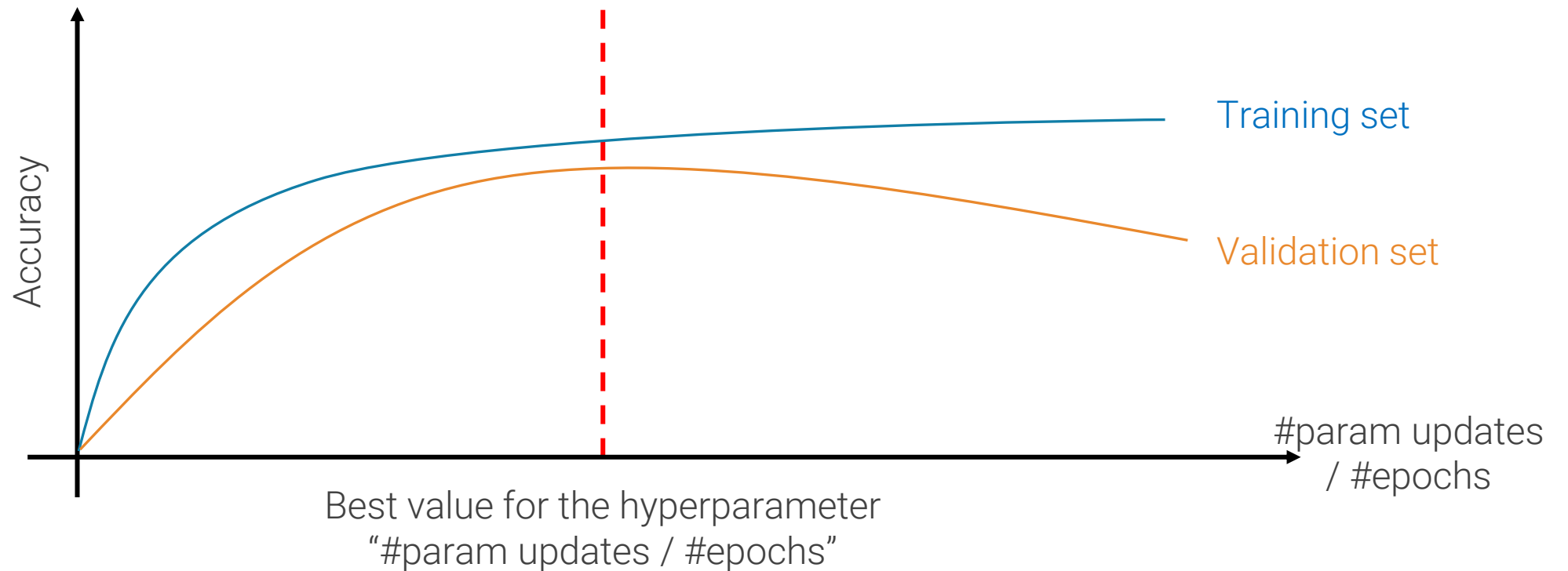
```
CLASS torch.optim.SGD(params, lr=<required parameter>, momentum=0, dampening=0,  
    λ weight_decay=0, nesterov=False, *, maximize=False, foreach=None) [SOURCE]
```

Stephen Jose Hanson and Lorien Y Pratt. "Comparing biases for minimal network construction with back-propagation.", NIPS 1988.



# Early stopping

Training time (i.e. number of parameter updates) is an hyperparameter controlling the effective capacity of the model. By using, at inference time, the best performing model on the validation set, we are effectively selecting the best value for this hyperparameter.



# Label smoothing

$s^{(i)}$  $\rightarrow$ 

$$-\log\left(\frac{\exp(s_{y^{(i)}})}{\sum_{k=1}^C \exp(s_k)}\right)$$

 $\leftarrow$  $\mathbb{I}(y^{(i)})$

$$LS(y^{(i)}, \epsilon)$$

0 (plane)	2,1
1 (car)	5,3
2 (bird)	7,6
...	0,2
	-1,3
	2,5
	3,5
	4,5
	5,5
	6,5

Cross-entropy loss has the one-hot encoding of the true label as target. Yet, it can not reach it: to have loss equal to 0, softmax of the correct label should be 1, hence  $s_{y^{(i)}} \rightarrow +\infty$  and  $s_{k, k \neq y^{(i)}} \rightarrow -\infty$   
Pushing scores in this way may cause overfitting

0 (plane)	0
1 (car)	0
2 (bird)	1
...	0
	0
	0
	0
	0
	0
	0

If we assume labels are corrupted by a small uniform noise  $\epsilon$ , we can mitigate the problem by using a **label smoothing** encoding.  
This also accounts for mislabeled examples.

0 (plane)	0.01	$= \epsilon / C$
1 (car)	0.01	$= \epsilon / C$
2 (bird)	0.91	$= 1 - \epsilon (C - 1) / C$
...	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$
	0.01	$= \epsilon / C$

# Label Smoothing in PyTorch v1.10 and above

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```
CLASS torch.nn.CrossEntropyLoss(weight=None, size_average=None, ignore_index=-100, reduce=None, reduction='mean', label_smoothing=0.0) \[SOURCE\]
```

Simply pass the correct class and set `label_smoothing > 0`

it is the epsilon of the previous slide

# Label Smoothing in PyTorch before v1.10

```
CLASS torch.nn.CrossEntropyLoss(weight: Optional[torch.Tensor] = None,  
size_average=None, ignore_index: int = -100, reduce=None, reduction: str =  
'mean') [SOURCE]
```

This criterion combines `nn.LogSoftmax()` and `nn.NLLLoss()` in one single class.

It is useful when training a classification problem with  $C$  classes. If provided, the optional argument `weight` should be a 1D *Tensor* assigning weight to each of the classes. This is particularly useful when you have an unbalanced training set.

The *input* is expected to contain raw, unnormalized scores for each class.

*input* has to be a *Tensor* of size either  $(\text{minibatch}, C)$  or  $(\text{minibatch}, C, d_1, d_2, \dots, d_K)$  with  $K \geq 1$  for the  $K$ -dimensional case (described later).

This criterion expects a class index in the range  $[0, C - 1]$  as the *target* for each value of a 1D tensor of size *minibatch*; if *ignore\_index* is specified, this criterion also accepts this class index (this index may not necessarily be in the class range).

CE loss expects only the value of the correct class for each example in the mini-batch, we can't pass the label smoothing encoding of  $y_i$  to it...

# Why cross-entropy?

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The name cross-entropy loss comes from the cross-entropy  $H(p, q)$ , a distance used in information theory between two probability distributions  $p$  and  $q$  over the same set of events, defined as

$$H(p, q) = -\mathbb{E}_p[\log q]$$

If  $p$  and  $q$  are **discrete** probability distributions, this becomes

$$H(p, q) = -\sum_{x \in \mathbb{X}} p(x) \log q(x)$$

In classification, the set of events  $\mathbb{X}$  is the set of the classes  $1, \dots, C$ , and a vector of  $C$  positive values which sum to one is a probability mass function over this set.

Therefore, we can think of the one-hot encoding of the true label as the distribution  $p$ , and the output of the model after the softmax, as the distribution  $q$ . Then, we recover the expression of the standard loss

$$H(\mathbb{I}(y^{(i)}), p_{model}(Y|x^{(i)}; \theta)) = -\sum_{k=1}^C \mathbb{I}(y^{(i)})_k \log p_{model}(Y = k|x^{(i)}; \theta) = -\log p_{model}(Y = y^{(i)}|x^{(i)}; \theta)$$

# KLDiv Loss

With the interpretation of the loss as the cross-entropy, it is easy to change the one-hot encoding with the label smoothing encoding of the true label.

$$H\left(\mathbf{LS}(y^{(i)}, \epsilon), p_{\text{model}}(Y|x^{(i)}; \theta)\right) = -\sum_{k=1}^C \mathbf{LS}(y^{(i)}, \epsilon)_k \log p_{\text{model}}(Y = k|x^{(i)}; \theta)$$

One last step: cross-entropy can be expressed in terms of other two information-theory quantities, the **Kullback-Leibler divergence**  $D_{KL}$  and the **entropy**  $H$ .

$$H(p, q) = H(p) + D_{KL}(p \parallel q)$$

Entropy of the labels  $H\left(\mathbf{LS}(y^{(i)}, \epsilon)\right)$  is constant, hence minimizing CE is equivalent to minimizing KL divergence from the model  $q$  to the labels  $p$  (**in this order**, divergence is NOT commutative).

```
CLASS torch.nn.KLDivLoss(size_average=None, reduce=None, reduction: str = 'mean',  
    log_target: bool = False)
```

**[SOURCE]** [🔗](#)

The **Kullback-Leibler divergence** Loss

target is  $p$ , i.e.  $\mathbf{LS}(y^{(i)}, \epsilon)$ ,  
input is  $\log(q)$ , i.e.  $\log\left(p_{\text{model}}(Y = k|x^{(i)}; \theta)\right)$

This criterion expects a *target Tensor* of the same size as the *input Tensor*.

# Dropout

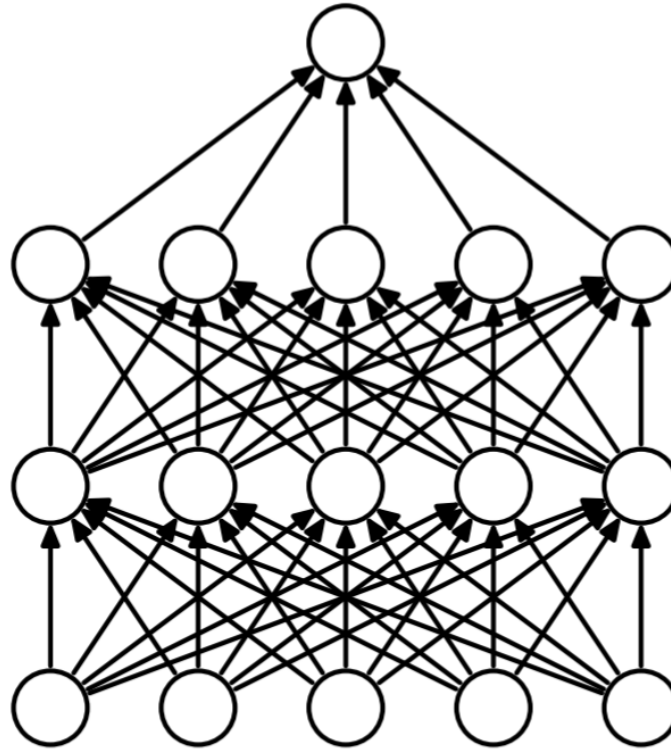
Arrows are weights and circles are activations

In each forward pass, **randomly set some activations to zero**

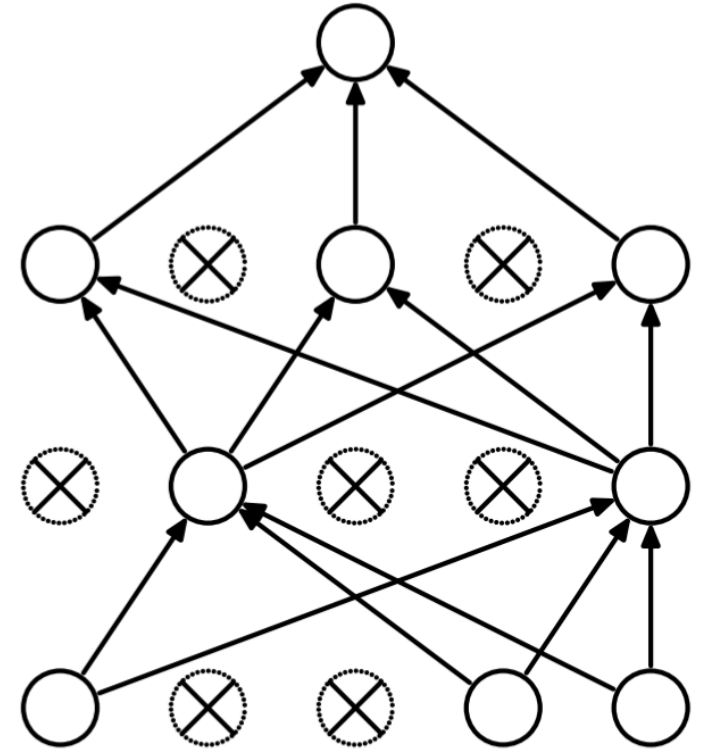
Probability of keeping an activation is a hyperparameter  $p$ , usually set to 0.5 and as high as 0.8 or 0.9.

Every time we train on a **new minibatch of examples**, we sample a **new binary mask** of active nodes.

We may apply it between any two layers, but never to the output activations!



Full network



Network used for one forward pass

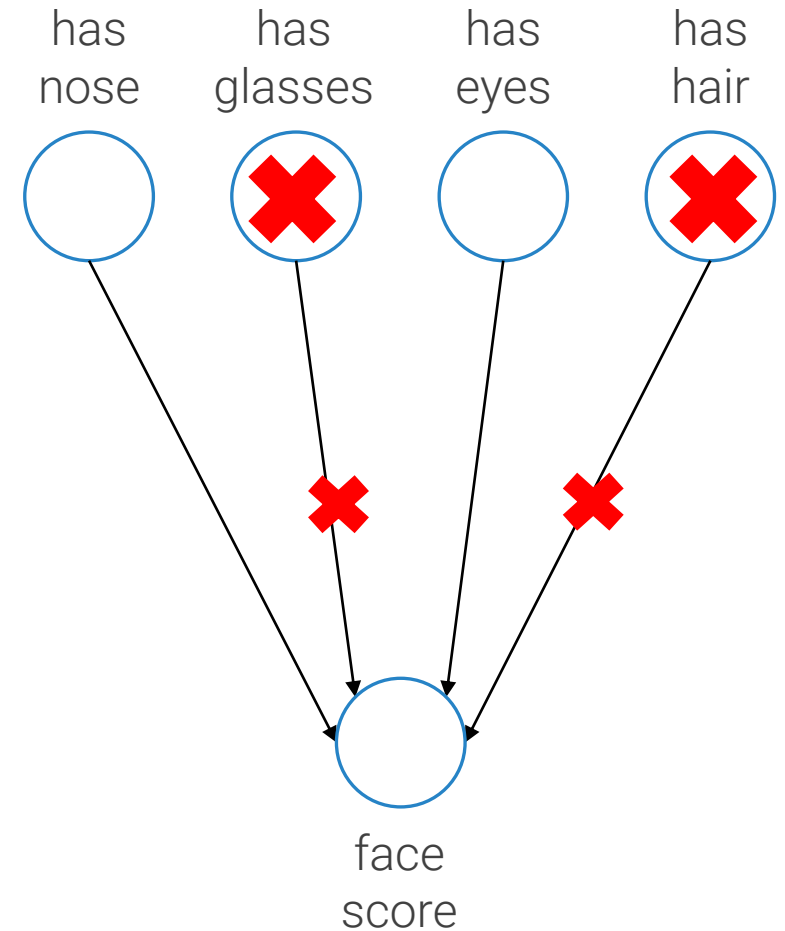
# Dropout as anti-conspiracy strategy

The idea of dropout was inspired to Hinton by his bank.

“I went to my bank. The tellers kept changing and I asked one of them why. He said he didn’t know but they got moved around a lot. I figured it must be because it would require cooperation between employees to successfully defraud the bank. This made me realize that randomly removing a different subset of neurons on each example would prevent conspiracies and thus reduce overfitting.”

Each hidden unit must be able to perform well regardless of which other hidden units are in the model, **it regularizes units to be useful in many context.**

As disruptive noise is applied to hidden units even in deep layers, makes it able to **destroy selective information** from the input, e.g. erases noses from faces and ask the network to recognize them anyway.





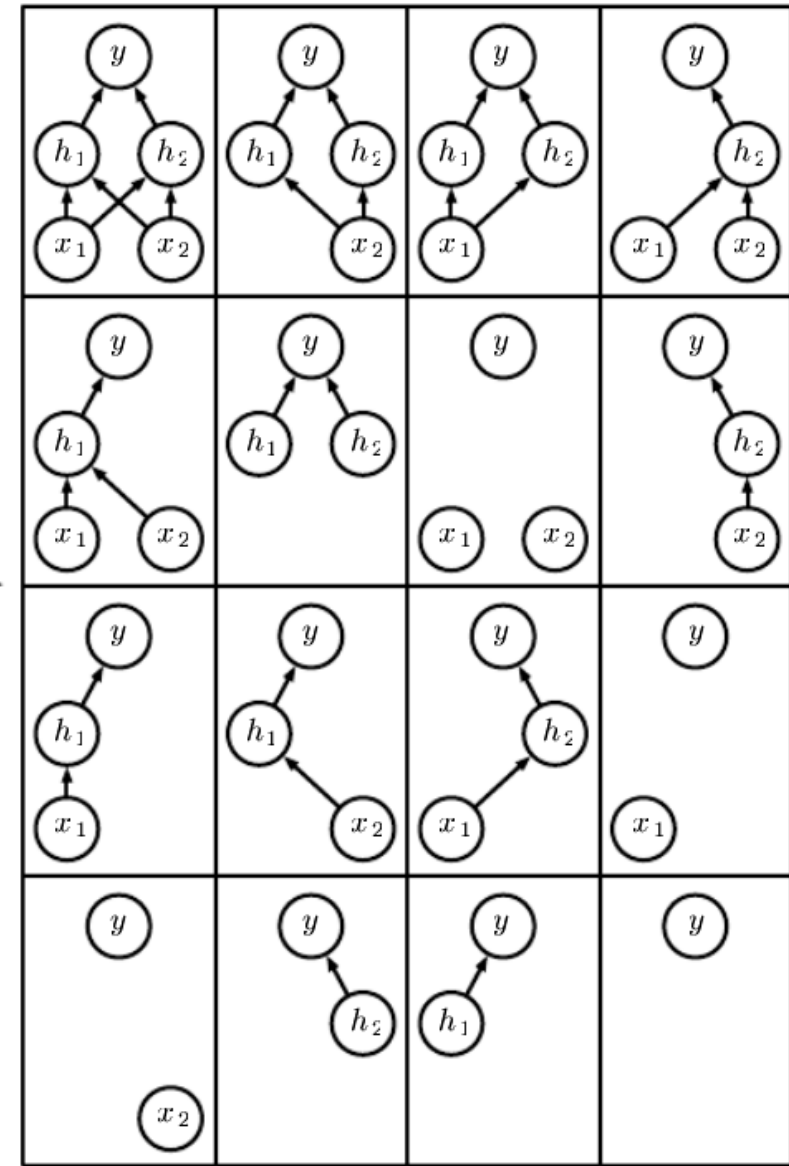
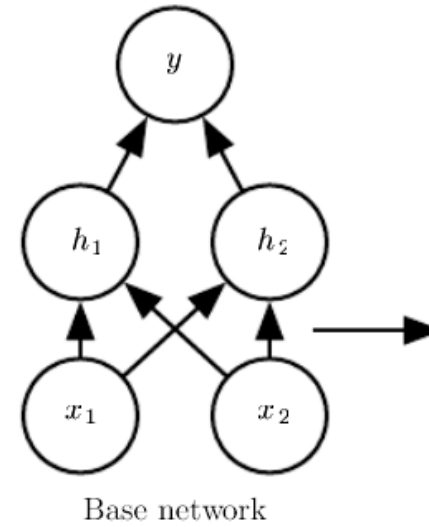
# Alternative interpretation: ensemble

We can think of dropout as training a large ensemble of models: each random binary mask is a new “model”.

Ensemble of models are a widely used technique to improve performance.

Key difference: “models” in dropout **share weights**

We can think of this ensemble as a special form of **bagging**, which we will study when discussing **Random Forests**



Ensemble of subnetworks

NOTE: configurations without connections between input and output highly unlikely in large models

# Dropout – Test time

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Dropout makes predictions at training time stochastic, but we want output at test time to be deterministic

$$scores = f(x; \theta, \textcolor{brown}{m})$$

 Random mask

Principled solution: “average out” stochasticity at test time

$$scores = f(x; \theta) = \mathbb{E}_{\textcolor{brown}{m}}[f(x; \theta, \textcolor{brown}{m})] = \sum_{\text{all masks } \textcolor{brown}{m}} p(\textcolor{brown}{m}) f(x; \theta, \textcolor{brown}{m})$$

We can approximate the sum, but everything is slowed down: e.g. for each test example  $x$ , pass it through the network several times sampling a different mask at each step (the more the better) and then average predictions.

# Dropout – Weight scaling

A faster approximation is provided by **weight scaling**

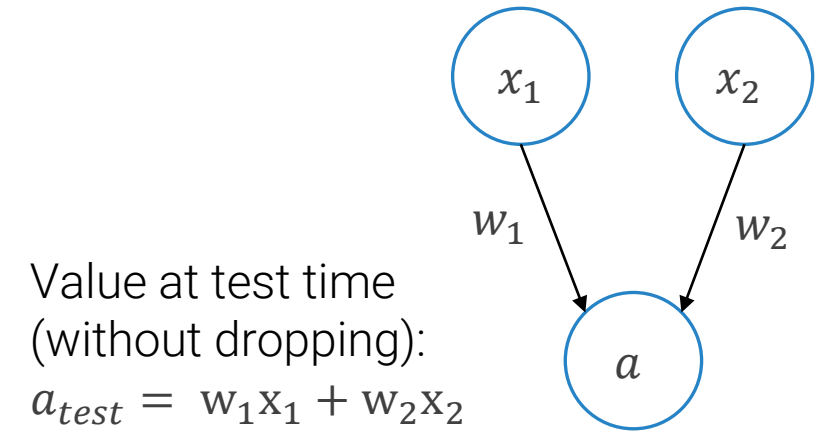
Consider an activation obtained by merging two **linear** neurons: in this case the summation can be solved explicitly, and it shows that at training time we obtain on average the **same activation without dropping rescaled by  $p$**

To make training and test time equal, we can either:

1. Rescale value at test time by  $p \rightarrow p a_{test} = \mathbb{E}_m[a_{train}]$
2. Rescale value at training time by  $\frac{1}{p} \rightarrow a_{test} = \mathbb{E}_m\left[\frac{a_{train}}{p}\right]$

The latter is referred to as **inverted dropout** and is preferred as it leaves test time unchanged.

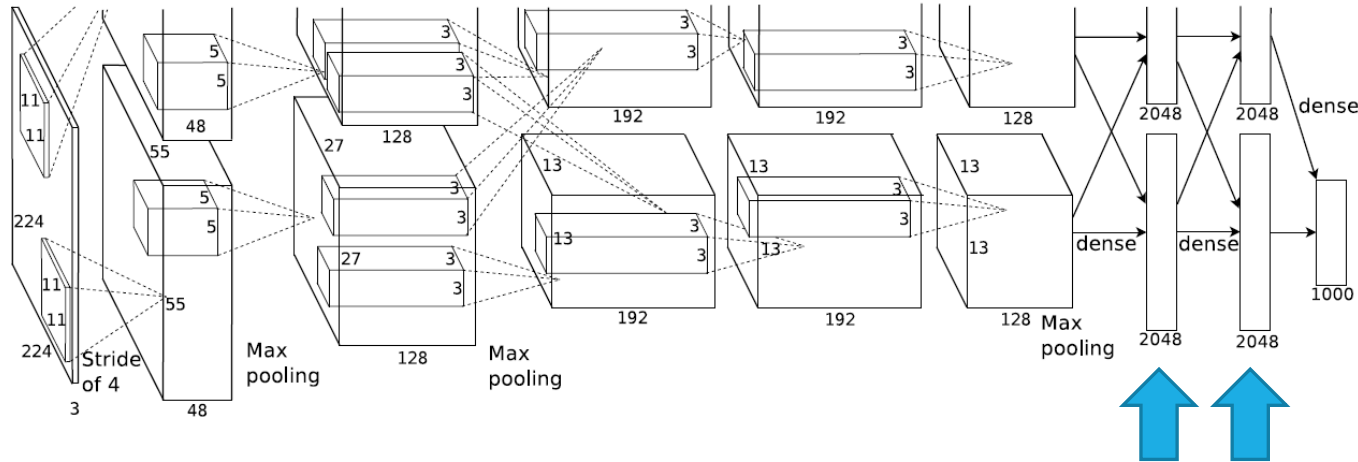
While this is **exact only for linear layers**, it is used as a fast approximation also in the presence of non-linearities.



Expected value at training time with  $p = 0.5$ :

$$\begin{aligned} \mathbb{E}_m[a_{train}] &= \frac{1}{4}(w_1 x_1 + w_2 x_2) + \frac{1}{4}(w_1 x_1 + 0) \\ &\quad + \frac{1}{4}(0 + w_2 x_2) + \frac{1}{4}(0 + 0) \\ &= \frac{1}{2}(w_1 x_1 + w_2 x_2) = p a_{test} \end{aligned}$$

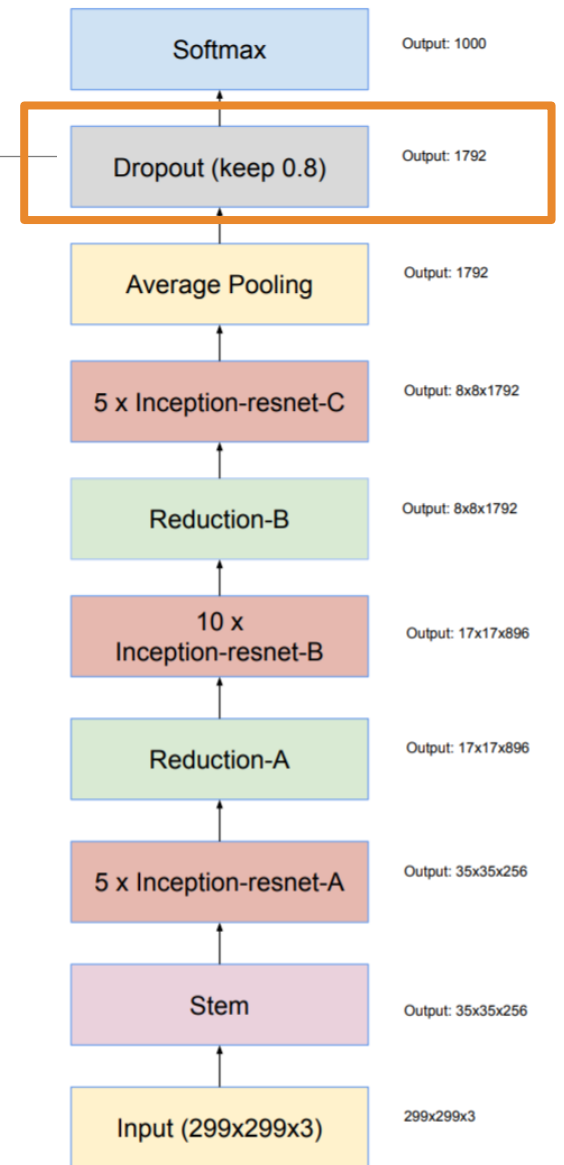
# Dropout: where



Usually applied to layers with many parameters, e.g. all but last FC layers in AlexNet, VGG, between GlobalAveragePooling and last fc layer in Inception-ResNet-v2,...

Not used in ResNets and variants, for its unclear interactions with BatchNorm.

EfficientNet uses it again, and scales  $p$  linearly from 0.2 to 0.5 going from B0 to B7.



# Regularization: a general template

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Training time: add randomness

Test time: average it out ( sometimes approximately

$$scores = f(x; \theta, \mathbf{m}) \quad f(x; \theta) = \sum_{\text{all masks } \mathbf{m}} p(\mathbf{m}) f(x; \theta, \mathbf{m})$$

Example: batch norm

At training time, each activation is randomly influenced by other samples in the mini-batch

$$\hat{a}_j^{(i)} = \frac{a_j^{(i)} - \mu_j}{\sqrt{v_j + \epsilon}}$$

$\mu_j$  and  $v_j$  **fixed to averages** seen at training time, BN becomes a deterministic linear operation

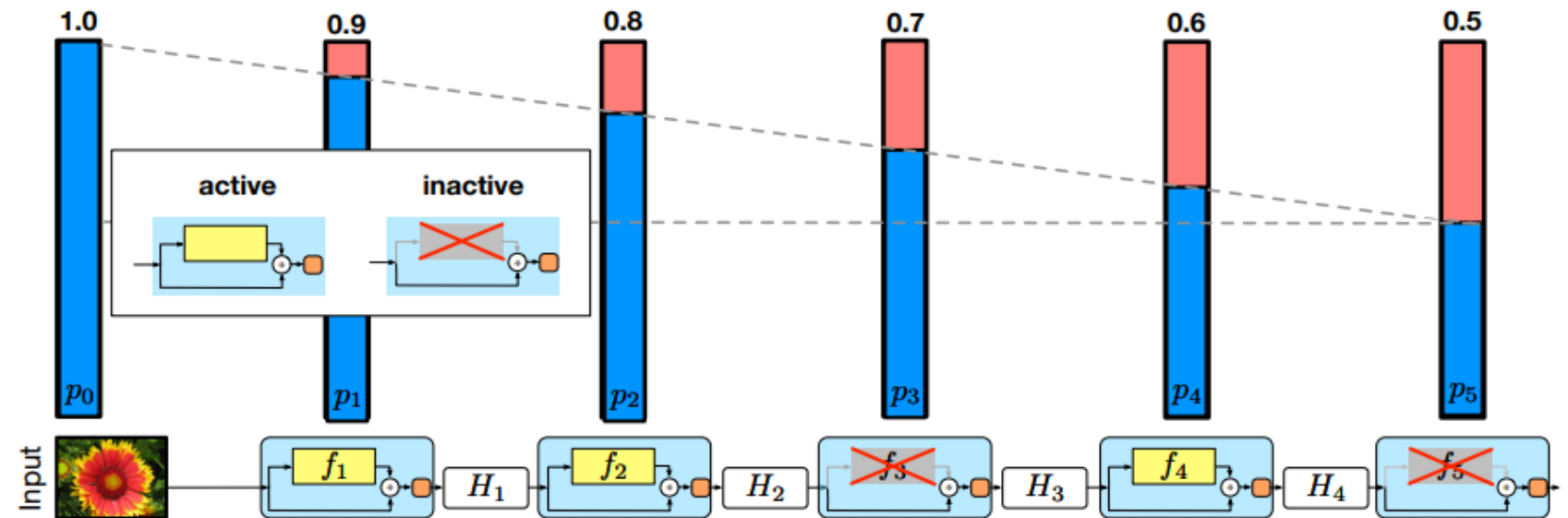
# Stochastic depth

Training time: **shrink depth**

For ResNet-like architectures, drop with probability  $1 - p_l$  the convolutional path of a ResNet-block, i.e. keep active only the residual identity path.

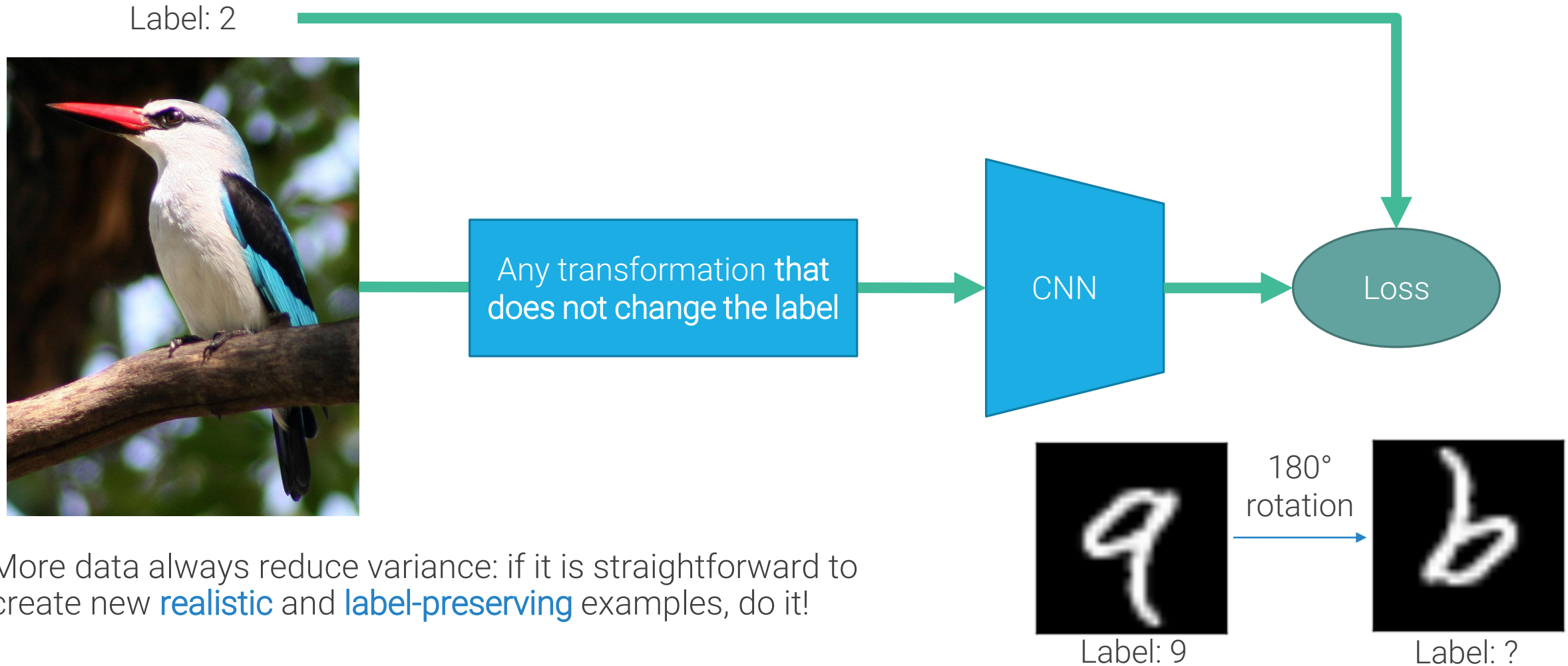
The survival probability decreases with depth

$$p_l = 1 - \frac{l}{L}(1 - p_L)$$



Test time: **unmodified network**

# Data augmentation

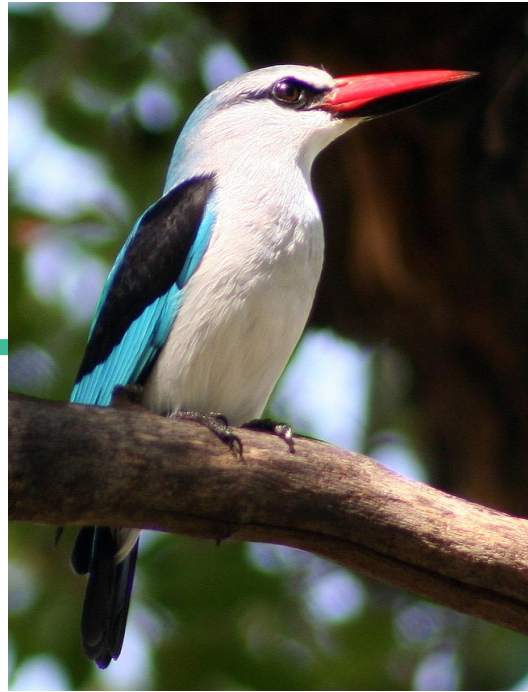


More data always reduce variance: if it is straightforward to create new **realistic** and **label-preserving** examples, do it!

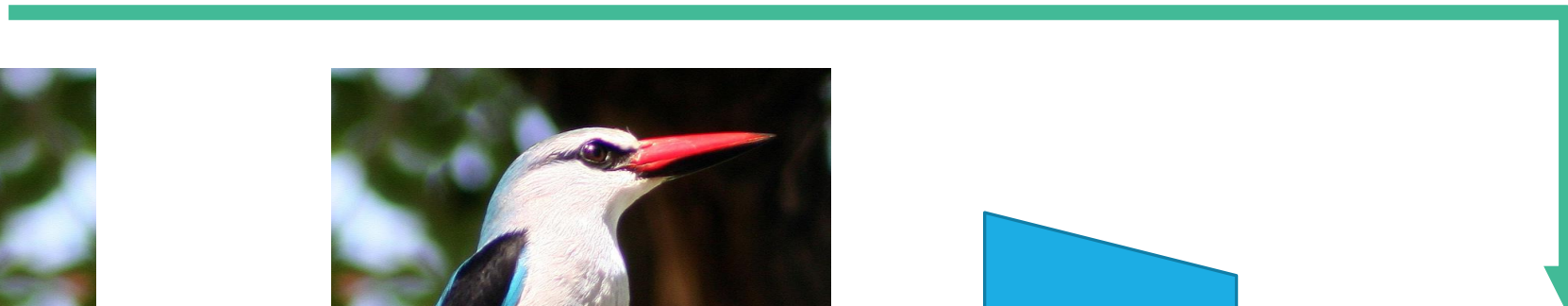
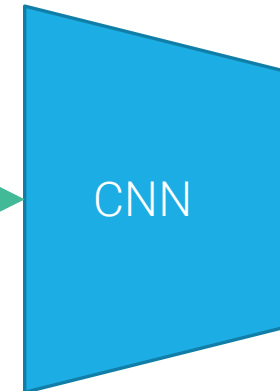


# Data augmentation: example

Label: 2



e.g. horizontal flip





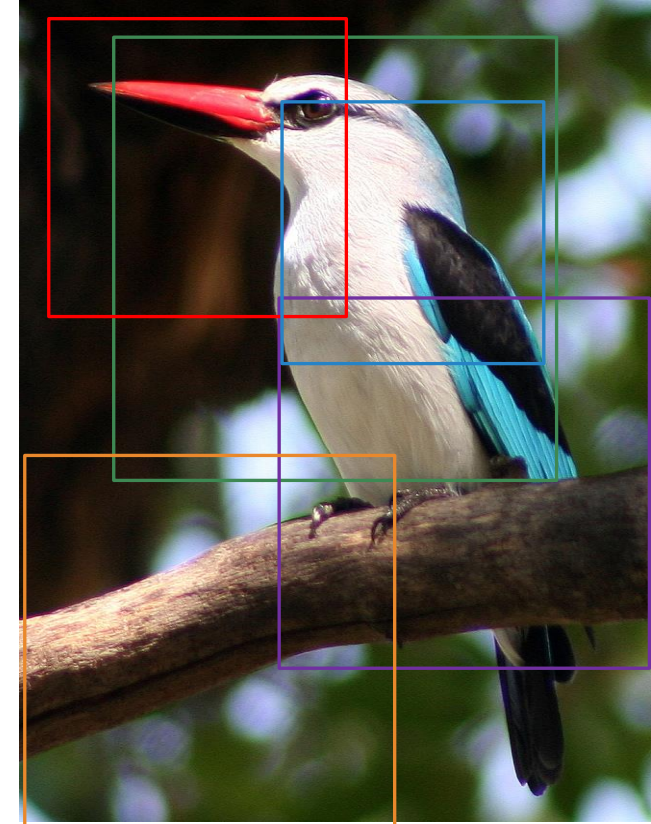
# Data augmentation: multi-scale training

Employed by ResNet, inherited from VGG

Training: **sample random crops / scales**

1. Pick random  **$S$**  in range  $[S_{\min}, S_{\max}] = [256, 480]$
2. Isotropically resize training image so that short side =  **$S$**
3. Sample random 224 x 224 **patch**

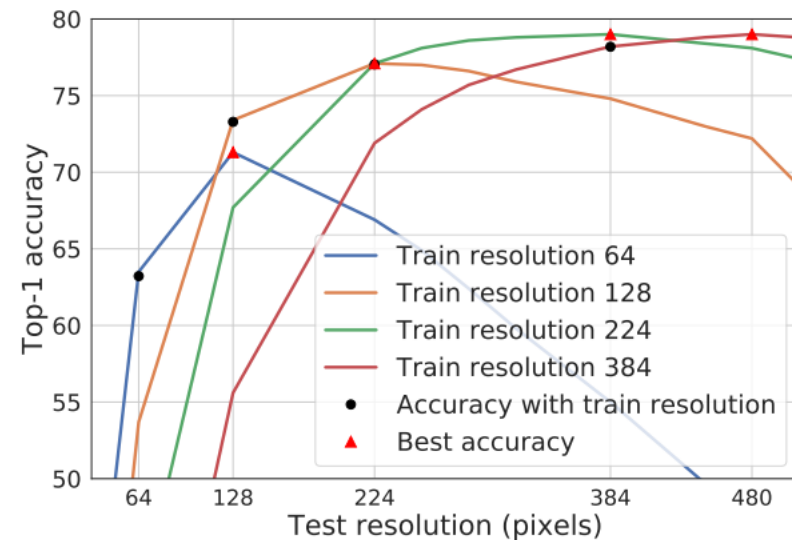
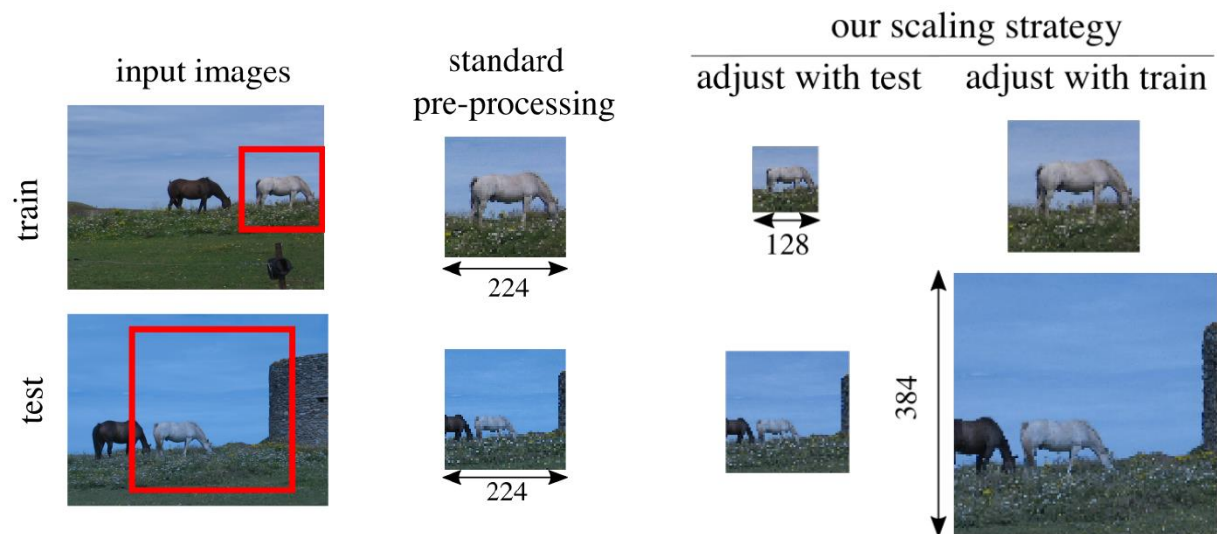
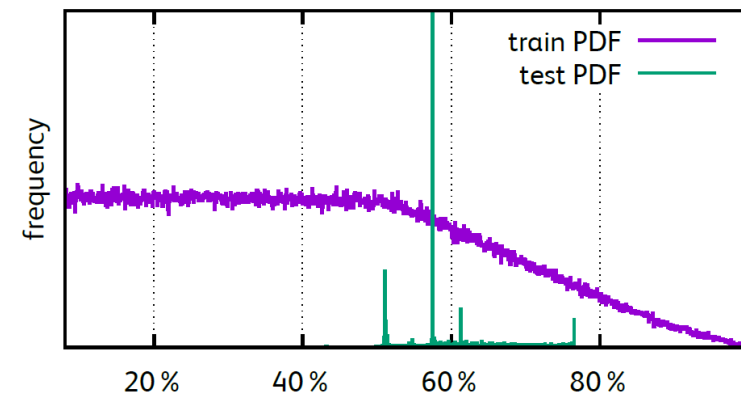
For smaller  $S$ , similar to 224, the crop will capture whole-image statistics, while for  $S \gg 224$  the crop will correspond to a small part of the image, containing a small object or an object part.



# FixRes

Data augmentations induce a significant **discrepancy between the size of the objects seen by the classifier at train and test time**: in fact, a lower train resolution improves the classification at test time!

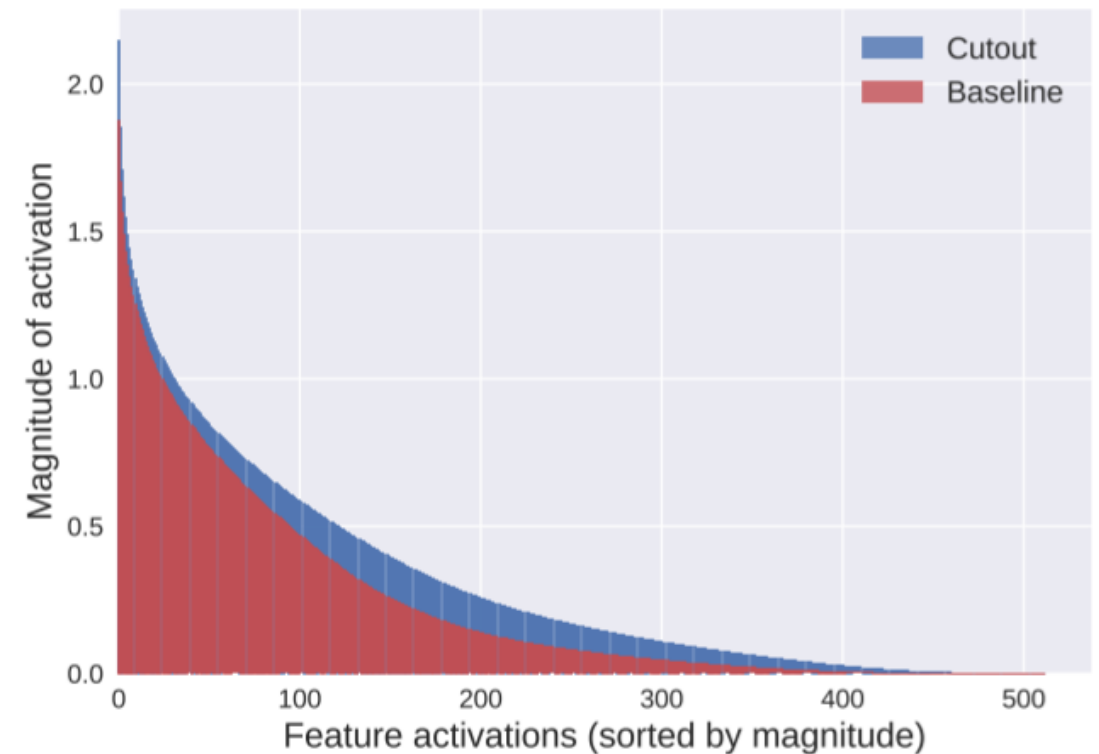
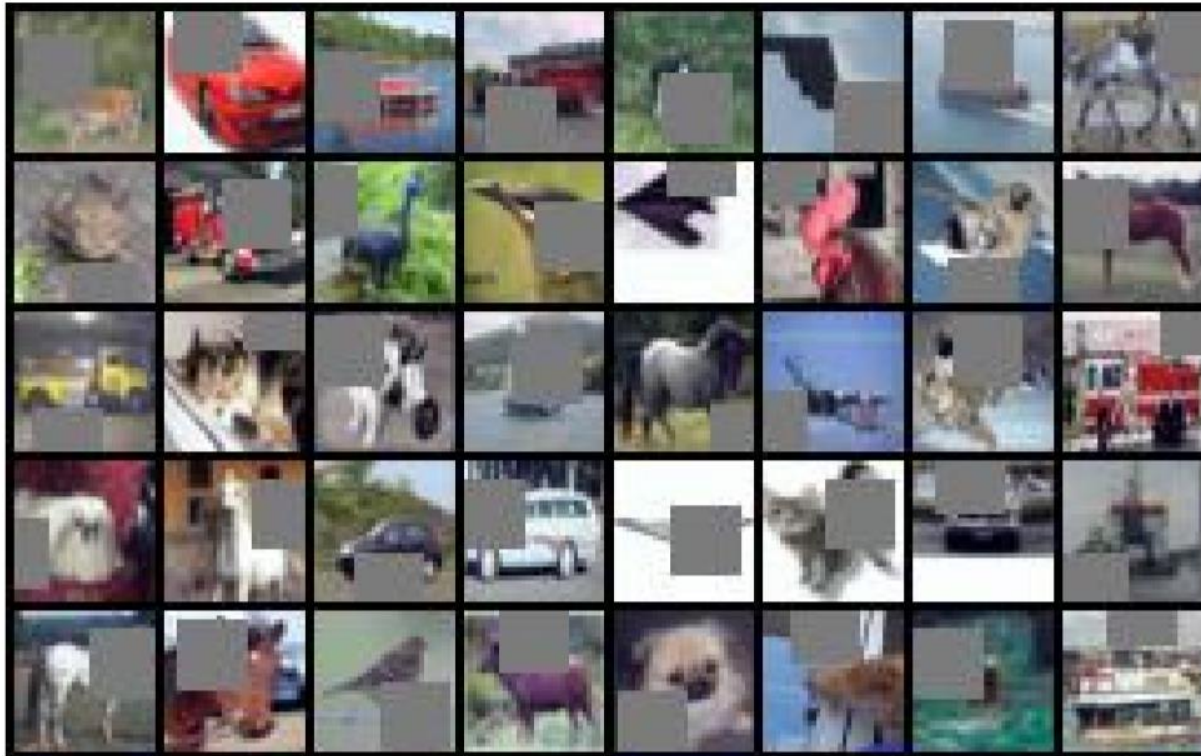
FixRes is a simple strategy to optimize the classifier performance, that fix the discrepancy by: (a) calibrating the object sizes by adjusting the crop size and (b) adjusting statistics of global average pooling by fine-tuning on larger crops the classifier and the last batch-norm layer before pooling.



Hugo Touvron et al., "Fixing the train-test resolution discrepancy", NeurIPS 2019.

# Cutout (Random Erasing)

Remove a random square region of the input image with 50% probability = “Dropout of the input space”. It forces the network to use a more diverse set of features, improving generalization.



Terrance DeVries and Graham W. Taylor, Improved Regularization of Convolutional Neural Networks with Cutout, 2017  
Zhun Zhong et al., “Random Erasing Data Augmentation”, AAAI 2020.



# Mixup

Training: **random blending of images**

Given two samples  $(x^{(i)}, y^{(i)})$ ,  $(x^{(j)}, y^{(j)})$

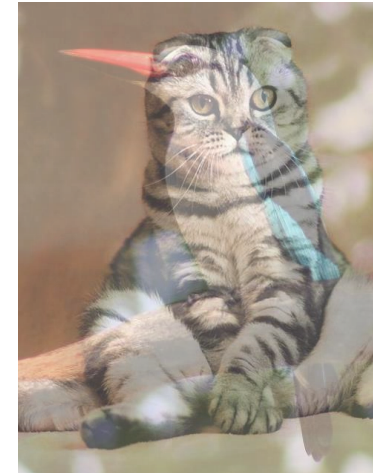
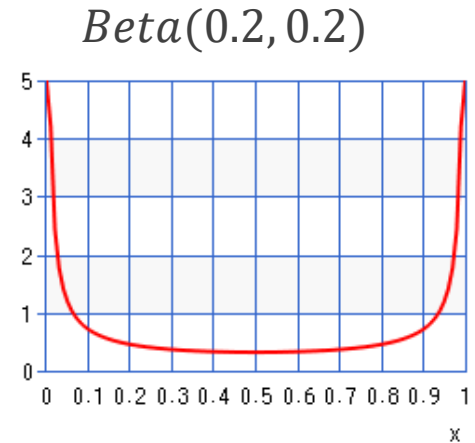
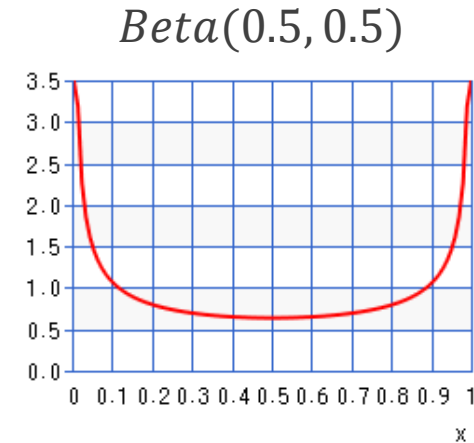
1. Sample **random blend probability**  $\lambda$  from a PDF  $Beta(\alpha, \alpha)$ ,  $\alpha \in (0, +\infty)$
2. Perform linear interpolation in the input space **and in the one-hot encoded label space** to define actual input sample and label

$$x = \lambda x^{(i)} + (1 - \lambda) x^{(j)}$$

$$y = \lambda \mathbb{I}(y^{(i)}) + (1 - \lambda) \mathbb{I}(y^{(j)})$$

Testing: **unmodified input**

Intuition: it forces the network to act linearly in between classes and have smoother transitions from one class decision boundary to another. Both effects help generalization.



0 (plane)	0
1 (cat)	0.6
2 (bird)	0.4
...	0
	0

# CutMix

Training: patches are cut and pasted among training images

Given two samples  $(x^{(i)}, y^{(i)})$ ,  $(x^{(j)}, y^{(j)})$  in a mini-batch

1. Sample random blend probability  $\lambda$  from a PDF  $Beta(\alpha, \alpha)$ ,  $\alpha \in (0, +\infty)$
2. Sample a rectangular region  $(r_x, r_y, r_h, r_w)$  whose aspect ratio is proportional to the original image and use it to define a binary mask  $M$  filled with 0 inside the region, 1 outside.

$$r_x \sim U(0, W) \quad r_y \sim U(0, H) \quad r_w = W\sqrt{1 - \lambda} \quad r_h = H\sqrt{1 - \lambda}$$

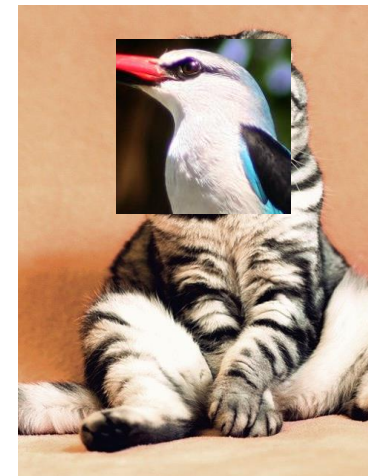
Note that the ratios of areas between region and image is  $\frac{r_w r_h}{WH} = 1 - \lambda$

3. Combine images according to mask and perform linear interpolation in the one-hot encoded label space to define the label

$$x = M \odot x^{(i)} + (1 - M) \odot x^{(j)}$$

$$y = \lambda \mathbb{I}(y^{(i)}) + (1 - \lambda) \mathbb{I}(y^{(j)})$$

Testing: unmodified input



0 (plane)	0
1 (cat)	0.8
2 (bird)	0.2
...	0
	0

# Random Hyper-parameters search

Grid search (log-linear):

$\forall lr \in 10^{\{-5, -4, -3, -2\}}$

$\forall wd \in 10^{\{-5, -4, -3, -2\}}$

train a model

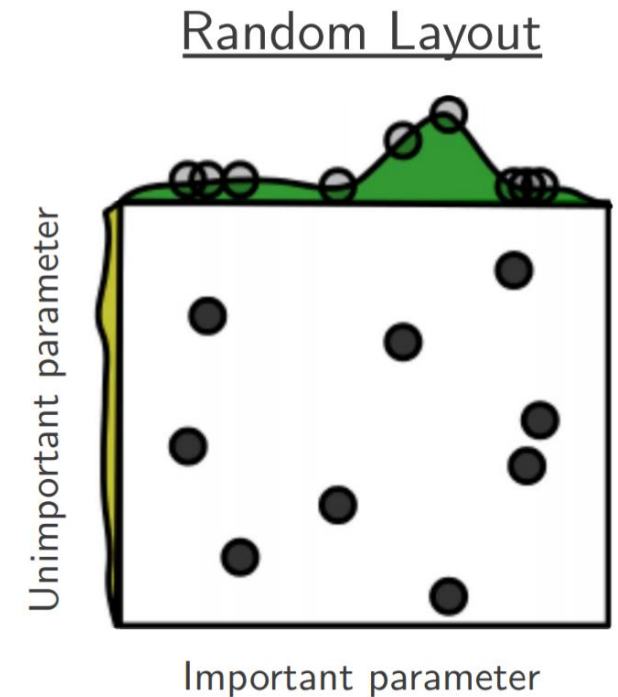
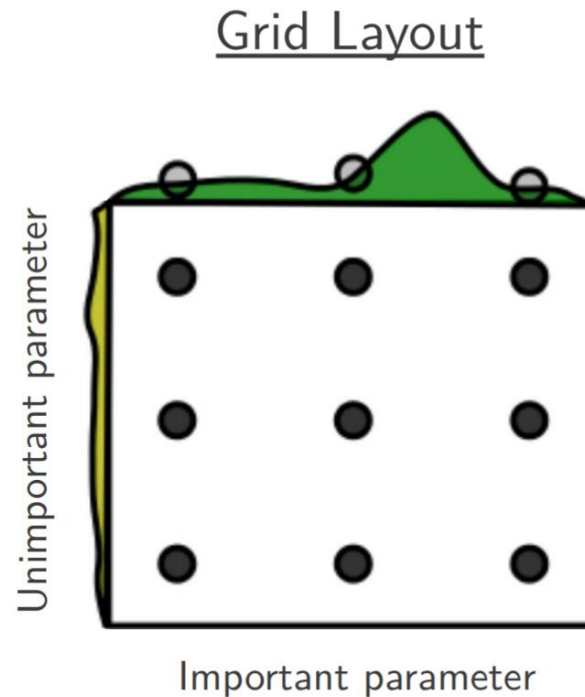
Random search: sample

$lr$  from  $10^U[-5, -2]$

$wd$  from  $10^U[-5, -2]$

and train a model.

It leads to a more efficient exploration of the space (but beware of the curse of dimensionality!)



# Test time good practice: ensembles

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1. Train multiple (randomly initialized) models on the same dataset.
2. Run each model over a test image, average the results (e.g. take average of logits, then take argmax)

This usually increases the overall performance by 1-2%: even if networks have similar error rates, they tend to make different mistakes.

Downside 1: we have to train a lot of networks from scratch

Downside 2: we have to run a lot of networks at test-time

# Exponential Moving Average (EMA)

To avoid building and running a costly ensemble, **we can average snapshots in weight space.**

**EMA** (sometimes called Polyak average): store and update with SGD the usual vector of parameters used at training time  $\theta$

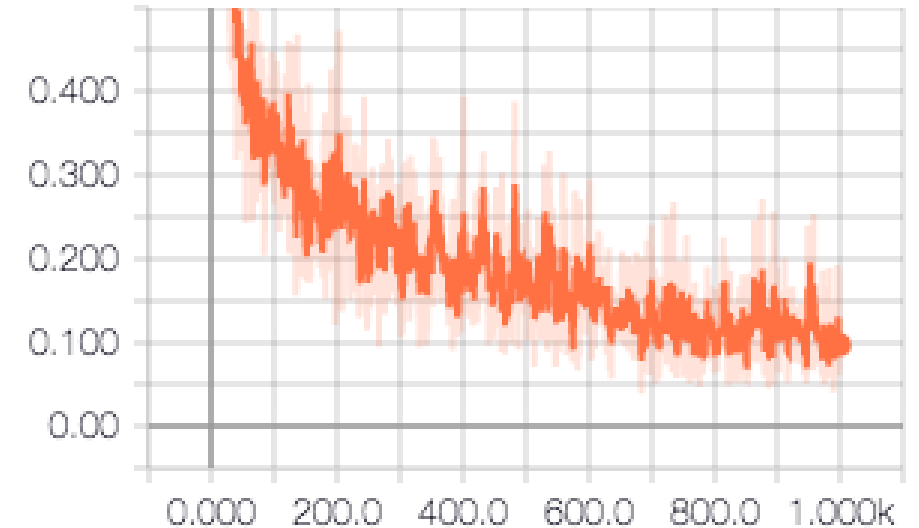
$$\theta^{(i+1)} = \theta^{(i)} - lr \nabla_{\theta} L(\theta; D^{(train)})$$

but also **store a vector of parameters to be used at test time, updated with an exponential moving average** every  $k$  step

$$\theta^{(test)} = (1 - \rho) \theta^{(i+1)} + \rho \theta^{(test)} \quad \rho \in [0,1]$$

( $\rho$  usually very large)

cross\_entropy\_1





# A “recipe” for training NNs

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“... suffering is a perfectly natural part of getting a neural network to work well, but **it can be mitigated by being thorough, defensive, paranoid, and obsessed with visualizations** of basically every possible thing. The qualities that in my experience correlate most strongly to success in deep learning are **patience and attention to detail.**”

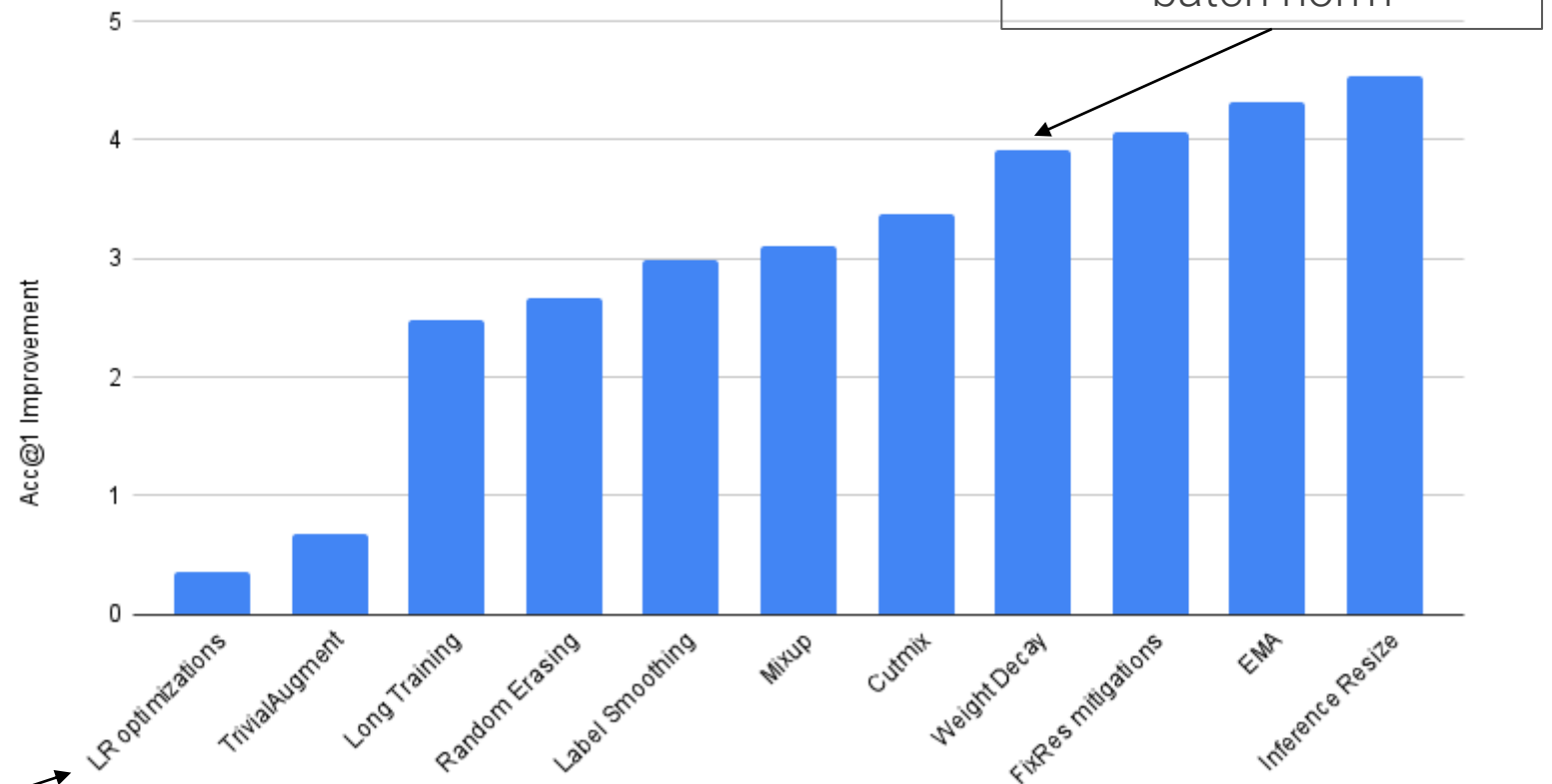
1. **Become one with the data:** collect statistics but also look at the data, understand what is relevant
2. **Set up the end-to-end training/evaluation skeleton + get dumb baselines:** check all the infrastructure code before training complex models, check init loss, **overfit a small dataset**, etc..
3. **Overfit:** reach low bias by starting with known models + Adam, then explore
4. **Regularize:** apply data augmentation, norm penalties, dropout, stochastic depth, cutmix, etc..
5. **Tune:** random search for better hyper parameters around what worked in 3-4, use LR schedules
6. **Test-time optimizations:** ensembles and/or distillation

# A case study: new weights in PyTorch

In 2021, PyTorch refreshed their pre-trained weights and API.

“... training models **is not a journey of monotonically increasing accuracies** and the process involves a lot of **backtracking**. To quantify the effect of each optimization, below we attempt to show-case **an idealized linear journey** of deriving the final recipe starting from the original recipe of TorchVision. We would like to clarify that this is an oversimplification of the actual path we followed and thus it should be taken with a grain of salt.”

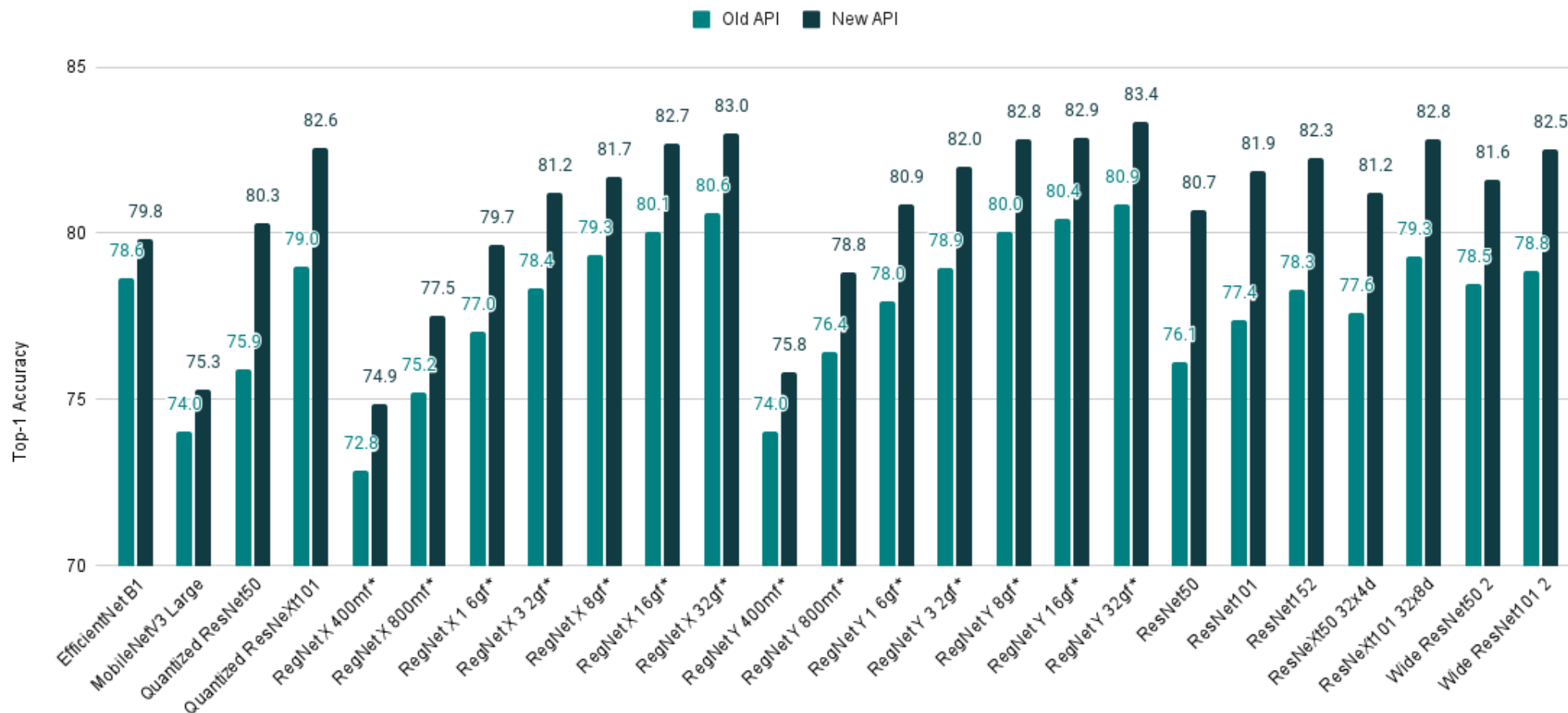
Cumulative Accuracy Improvements for ResNet50



lr value + warmup + cosine schedule

<https://pytorch.org/blog/how-to-train-state-of-the-art-models-using-torchvision-latest-primitives/>

# Training recipe matters



<https://pytorch.org/blog/introducing-torchvision-new-multi-weight-support-api/>